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(54) Title: POLYMORPHISMS AND NEW GENES IN THE REGION OF THE HUMAN HEMOCHROMATOSIS GENE

(57) Abstract

Polymorphic sites in the region surrounding the HFE gene are provided. These polymorphisms are useful as surrogate markers in diagnostic assays for hemochromatosis. Additionally, a fine structure map of the 1 megabase region surrounding the HFE gene is provided. along with 235 kb of DNA sequence and 8 loci corresponding to candidate genes within the 1 megabase region, and in the purification of related proteins.

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Polymorphisms and New Genes in the Region of the Human Hemochromatosis Gene

BACKGROUND OF THE INVENTION

Hereditary hemochromatosis (HH) is an inherited disorder of iron metabolism wherein the body accumulates excess iron. In symptomatic individuals, this excess iron leads to deleterious effects by being deposited in a variety of organs leading to their failure, and resulting in cirrhosis, diabetes, sterility, and other serious illnesses. The gene which is defective in this disease was disclosed in copending U.S.S.N. 08/652,265.

Fine structure mapping of the region to which the gene responsible for HH, HFE (denoted HH or HFE in some publications), was mapped makes possible the identification of candidate sequences comprising the HFE gene, along with structural elements for regulation and expression and neighboring genes.

A variety of techniques is available for fine structure mapping, including direct cDNA selection, exon-trapping, and genomic sample sequencing. The direct selection approach (Lovett et al. Proc. Natl. Acad. Sci. U.S.A. 88:9628-9623 (1991)) involves the hybridization of cDNA fragments to genomic DNA. This technique is extremely sensitive and capable of isolating portions of rare transcripts. Exon-trapping (Church et al. Nature Genetics 6:98-105 (1994)) recovers spliced introns from in vivo expressed genomic DNA clones and produces candidate exons without requiring any prior knowledge of the target's gene expression. High-throughput genomic DNA sequencing with comparison of the sequence data to databases of expressed sequences has also been used, such as in the positional cloning of the Werner syndrome gene. (Yu et al. Science 277:258-262 (1996)) and in cloning by homology of the second Alzheimer's disease gene on chromosome 1 (Levy-Lahad et al. Science 269:973-977 (1995)).

HH is typically inherited as a recessive trait; in the current state of knowledge. homozygotes carrying two defective copies of the gene are most frequently affected by the disease. In addition, heterozygotes for the HFE gene are more susceptible to sporadic porphyna cutanea tarda and potentially other disorders (Roberts et al., <u>Lancet 349:321-323 (1997)</u>. It is estimated that approximately 10-15% of Caucasians carry one copy of the HFE gene mutation and that there are about one million homozygotes in the United States. HH, thus, represents one of the most common genetic disease mutations in Caucasian individuals. Although ultimately HH produces debilitating symptoms, the majority of homozygotes and heterozygotes have not been diagnosed.

The need for such diagnostics is documented, for example, in Barton, J.C. et al. Nature Medicine 2:394-395 (1996); Finch, C.A. West J Med 153:323-325 (1990); McCusick, V. Mendelian Inheritance in Man pp. 1882-1887, 11th ed., (Johns Hopkins University Press, Baltimore (1994)); Report of a Joint World Health Organization/Hemochromatosis Foundation/French Hemochromatosis Association Meeting on the Prevention and Control of Hemochromatosis (1993); Edwards, C.Q. et al. New Engl J Med 328:1616-1620 (1993); Bacon, B.R. New Engl J Med 326:126-

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A furth it aspect of the invention is an isolated nucleic acid sequence comprising a	
nucleic acid sequence substantially identical to BTF5.	
A further aspect of the invention is an isolated nucleic acid sequence comprising a	
nucleic acid sequence substantially identical to BTF4.	
A further aspect of the invention is an isolated nucleic acid sequence comprising a	32
nucleic acid sequence substantially identical to BTF3.	
A further aspect of the invention is an isolated nucleic acid sequence comprising a	
nucleic acid sequence substantially identical to BTF2.	
s gnisher aspect of the invention is an isolated nucleic acid sequence comprising a	
acid sequence substantially identical to BTF1.	30
One aspect of the invention is an isolated nucleic acid sequence comprising a nucleic	
designation ATCC CRL-12371.	
Another aspect of the invention is a culture of lymphoblastoid cells having the	
genome of the individual.	
and the presence of the genotype indicates the likely presence of the HFE gene mutation in the	52
allele of Table 1 indicates the likely absence of the HFE gene mutation in the genome of the individual	
wherein, as a result, the absence of a genotype defined by a polymorphic	
defined by a polymorphic allele of Table 1,	÷
assessing the DNA or RNA for the presence or absence of a genotype	
providing DNA or RNA from the individual; and	50
the common hereditary hemochromatosis (HFE) gene mutation in an individual comprising:	
Another aspect of the invention is a method to determine the presence or absence of	
indicates the likely presence of the HFE gene mutation in the genome of the individual.	
absence of the HFE gene mutation in the genome of the individual and the presence of the haplotype	
wherein, as a result, the absence of a haplotype of Table 1 indicates the likely	۶ı
Table 1,	
assessing the DNA or RNA for the presence or absence of a haplotype of	
providing DNA or RNA from the individual; and	
the common hereditary hemochromatosis (HFE) gene mutation in an individual comprising:	
Another aspect of the invention is a method to determine the presence or absence of	10
DNA molecule comprises at least one polymorphic site of Table 1.	
100 consecutive bases to about 235 kb substantially identical to the sequence of Figure 9, wherein the	
Another aspect of the invention is an isolated nucleic acid molecule comprising about	
of Figure 9 or its complement for amplification of a polymorphic site of Table 1.	
Another aspect of the invention is an oligonucleotide pair selected from the sequence	S
least 8 to about 100 consecutive bases includes at least one polymorphic site of Table 1.	
consecutive bases from the sequence of Figure 9, or the complement of the sequence, wherein the at	
One aspect of the invention is an oligonucleotide comprising at least 8 to about 100	
SOMMAN OF THE INVENTION	

nucleic acid sequence substantially identical to NPT3.

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alignment represent amino acids conserved in all 6 proteins; the "dots" represent conserved amino acids substitutions. Boxed are the regions within the proteins which correspond to three conserved motifs: 1) the B-G domain, 2) the transmembrane domain (TM), and 3) the B30-2 exon domain.

Figure 4, panel (A) depicts a Northern blot analysis of representative members of the two groups of BTF proteins, BTF1 and BTF5. BTF1 hybridized to all tissues on the blot as a major transcript at 2.9 kb and a minor one at 5.0 kb. BTF5 hybridized to several transcripts ranging between 4.0 and 3.1 kb and as a similar expression profile to BTF1. Autoradiography was for 24 hours. The β-actin hybridization demonstrated the variation in ploy (A)+ RNA between the lanes. Autoradiography was for 1 hour. In panel (B), RT-PCR analysis demonstrated that the expression of both genes was widespread. Included in the (+) lane are cDNA 21 and 44 as positive controls; the (-) lane represents the no-DNA control. Amplification using primers for the RFP gene (Isomura *et al.* <u>Nucleic Acid Res.</u> 20:5305-5310 (1992)) controlled for the integrity of the cDNA. All first strand cDNAs were checked for contaminating genomic DNA amplification by carrying out an identical experiment excluding the reverse transcriptase. In all cases, no amplification was obtained (data not shown).

Figure 5(A) depicts an alignment of the predicted amino acid sequence of the RoRet gene to the 52 kD Ro/SSA auto-antigen protein. The asterisks under the alignment represent conserved amino acids; the "dots" represent conserved amino acids substitutions. The putative DNA binding cysteine-rich domain and the B30-2 exon domain are boxed. Figure 5(B) depicts an alignment of the predicted amino acid sequence of the two novel putative sodium phosphate transport proteins to that of the NPT1.

Figure 6, panel (A) depicts a Northern blot analysis of the RoRet gene. The RoRet cDNA hybridized to 4 different transcripts, ranging from 7.1 kb to 2.2 kb. Autoradiography was performed for 4 days. The re-hybridization of the blot with a β-actin probe showed the variation in poly (A)+ RNA between the lanes. Autoradiography was for 1 hour. Panel (B) depicts RT-PCR analysis of the RoRet gene. Included in the (+) lane was a cDNA 27 positive control. Weak amplification of the correct size was observed in the small intestine, kidney and liver. The other tissues were negative as was the no DNA control lane (-). The RFP primers demonstrated the integrity of the cDNA. Panel (C) depicts Northern blot analysis of NPT3 and NPT4. NPT3 was expressed at high abundance in the heart and muscle as a single 7.2 kb transcript. Lesser amounts were found in the other tissues. The expression pattern of NPT4 was more restricted, being found only in the liver and kidney as a smear of transcripts ranging from 2.6 to 1.7 kb. Panel (D) depicts RT-PCR analysis of the NPT3 and NPT4 genes. Included in the (+) lane were the respective cDNA22E and 22B positive controls. The NPT3 gene was expressed as the proper size PCR fragment in kidney, liver, spleen and testis. A smaller fragment was detected in all tissues with the exception of the liver. The no DNA control lane (-) was negative. NPT4 was expressed as the proper size fragment in the small intestine, kidney, liver and testis. Larger and smaller size fragments were found in all other tissues with the exception of the brain. For both genes these different size fragments may indicate alternative splice events. The no DNA control lane (-) was negative. The RFP primers demonstrated the integrity of the cDNA.

Figure 7 depicts the sequences of cDNA 21 (BTF1), cDNA 29 (BTF3), cDNA 23 (BTF4), cDNA 44 (BTF5), cDNA 32 (BTF2), cDNA 27 (RoRet), cDNA 22B (NPT3), cDNA22E (NPT4).

Interscience, New York (1987). Cuttent Protocols in Molecular Biology, F. Ausubel et al., ed. Greene Publishing and Wiley-Molecular Cloning: a Laboratory Manual (2nd ed.), Vols. 1-3, Cold Spring Harbor Laboratory, (1989) or discussions of nucleic acid probe design and annealing conditions, see, for example, Sambrook et al.,

provide codon preference in a specific host cell. includes the degenerate codons of the native sequence or sequences which may be introduced to length sequences derived from the full length protein. It being further understood that the sequence The nucleic acid sequences include both the full length nucleic acid sequences as well as non-full strand sequence that is transcribed into RNA and the RNA sequence that is translated into protein. the expression of a specific protein or peptide. The nucleic acid sequences include both the DNA The phrase "nucleic acid sequence encoding" refers to a nucleic acid which directs

The phrase "expression cassette", refers to nucleotide sequences which are capable lack at least one protein or nucleic acid normally associated with the nucleic acid in a host cell. The phrase "isolated" or "substantially pure" refers to nucleic acid preparations that

The term "operably linked" as used herein refers to linkage of a promoter upstream necessary or helpful in effecting expression may also be used as described herein. include at least promoters and optionally, transcription termination signals. Additional factors of affecting expression of a structural gene in hosts compatible with such sequences. Such cassettes

The term "vector", refers to viral expression systems, autonomous self-replicating from a DNA sequence such that the promoter mediates transcription of the DNA sequence.

replicated by the cells during mitosis as an autonomous structure, or is incorporated within the host's chromosome(s). Where a vector is being maintained by a host cell, the vector may either be stably both extrachromosomal circular DNA and DNA that has been incorporated into the host recombinant microorganism or cell culture is described as hosting an "expression vector," this includes circular DNA (plasmids), and includes both expression and nonexpression plasmids. Where a

The term "plasmid" refers to an autonomous circular DMA molecule capable of spliced from the mRNA transcript, along with variants resulting from atternative splice sites. enhancers, and termination regions. The term further includes all introns and other DNA sequences "gene" is intended to include not only coding sequences but also regulatory regions such as promoters, sequence variants wherein such afterations do not affect the function of the gene product. The term encodes a polypeptide. This definition includes various sequence polymorphisms, mutations, and/or The term "gene" as used herein is intended to refer to a nucleic acid sequence which

replicated by the cells during mitosis as an autonomous structure or is incorporated within the host's chromosome(s). Where a plasmid is being maintained by a host cell, the plasmid is either being stably extrachromosomal circular DNA molecules and DNA that has been incorporated into the host microorganism or cell culture is described as hosting an "expression plasmid", this includes both replication in a cell, and includes both the expression and nonexpression types. Where a recombinant

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The phrase "substantially purified" or "isolated" when referring to a peptide or protein, means a chemical composition which is essentially free of other cellular components. It is preferably in a homogeneous state although it can be in either a dry or aqueous solution. Purity and homogeneity are typically determined using analytical chemistry techniques such as polyacrylamide gel electrophoresis or high performance liquid chromatography. A protein which is the predominant species present in a preparation is substantially purified. Generally, a substantially purified or isolated protein will comprise more than 80% of all macromolecular species present in the preparation. Preferably, the protein is purified to represent greater than 90% of all macromolecular species present. More preferably the protein is purified to greater than 95%, and most preferably the protein is purified to essential homogeneity, wherein other macromolecular species are not detected by conventional techniques.

The phrase "specifically binds to an antibody" or "specifically immunoreactive with", when referring to a protein or peptide, refers to a binding reaction which is determinative of the presence of the protein in the presence of a heterogeneous population of proteins and other biologies. Thus, under designated immunoassay conditions, the specified antibodies bind to a particular protein and do not bind in a significant amount to other proteins present in the sample. Specific binding to an antibody under such conditions may require an antibody that is selected for its specificity for a particular protein. A variety of immunoassay formats may be used to select antibodies specifically immunoreactive with a particular protein. For example, solid-phase ELISA immunoassays are routinely used to select monoclonal antibodies specifically immunoreactive with a protein. See Harlow and Lane (1988) Antibodies, a Laboratory Manual, Cold Spring Harbor Publications, New York, for a description of immunoassay formats and conditions that can be used to determine specific immunoreactivity.

As used herein, "EST" or "Expressed Sequence Tag " refers to a partial DNA or cDNA sequence of about 150 to 500, more preferably about 300, sequential nucleotides of a longer sequence obtained from a genomic or cDNA library prepared from a selected cell, cell type, tissue or tissue type, or organisms which longer sequence corresponds to an mRNA or a gene found in that library. An EST is generally DNA. One or more libraries made from a single tissue type typically provide at least 3000 different (i.e. unique) EST's and potentially the full complement of all possible EST's representing all possible cDNAs, e.g., 50,000 - 100,000 in an animal such as a human. (See, for example, Adams et al. Science 252:1651-1656 (1991)).

"Stringent" as used herein refers to hybridization and wash conditions of 50% formamide at 42°C. Other stringent hybridization conditions may also be selected. Generally, stringent conditions are selected to be about 5° C lower than the thermal melting point (Tm) for the specific sequence at a defined ionic strength and pH. The Tm is the temperature (under defined ionic strength and pH) at which 50% of the target sequence hybridizes to a perfectly matched probe. Typically, stringent conditions will be those in which the salt concentration is at least about 0.02 molar at pH 7 and the temperature is at least about 60°C. As other factors may significantly affect the stringency of hybridization, including, among others, base composition and size of the complementary strands, the presence of organic solvents and the extent of base mismatching, the combination of parameters is more important than the absolute measure of any one.

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in the identification and isolation of further members of the gene family. Nucleic acid sequences substantially identically to the NPT1-like sequences and the proteins encoded by them are also included in the scope of this invention.

C. Polymombic Markers

C. <u>Polymorphic Markers</u>

These polymorphisms are listed in Table 1. As described below, these polymorphisms were identified by comparison of the DNA sequence of an affected individual homozygous for the common ancestral HH mutation with that of an unaffected individual disclosed in copending U.S. 08/724,394.

10 Table 1. Polymorphic Sites in the HH Region

ר	19721	D-A	38526	C-1
	16211	T-A	11478	9-Y
	61091	T DEL	58625	9-∀
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.]	12904-12919	ASTITSTASTSAASS	30725	Ø-9
≔ (78821	9-A	36958	Q-A
1	12264	0-1	32179	A-C
.1	86611	T-0	34434	T-D
SÞ	71441-31441	SNI A	330S6	130 T
.]	99461	A-Ð	33017	9-1
.]	13304	A-5	35226-32559	TGTG DEL
.]	12173-12175	130 TTT	35040	9-0
.]	99611	. D-T	31749	T-O
40	47811	9-A	31280	T-O
]	11224	C-A	91059	∀-T
.]	10955	9-A	30400	
	81601	9-0	77108	
	10828	D-A	30008	2-1
-	10214	T-O	S286S	D-A
	10027	A-9	78762	Ð-T
	5823	Y-9	Z9454-29457	130 TTT
	SII6	∀ -9	29100	V-9
	9478	D-T	28132	Q-P
	7933	D-T	19872	T DEL
4	2797	G DEL	27838	1-9
		DEL		!
	8227-2427	DETTABODADADADT	7693	<u>⊃-T</u>
<u>.</u>	7273	A-9	76667	Υ-Ο
-	9817	2.1	52175	ე- <u>១</u>
	8699	D-T	24143	A-D A-T
	<u> </u>	A DEL	<u> </u>	A-⊃
⊣	6231	A-Ð	22786	D-A A-2
	<u></u>	9-0	55593	2-4
	1109	Y-5	21837	2-4
	6888	T-O O-T	21117	T-A
	1699		20841	T-A
	4925-4928	C-G	20463	C-A
	3829	9-7	20366-20367	2NI A
	3767 2662-2663	130 TT	20089	1-0
→	148	0-1	67661	1-0
	35-36	AC DEL	99261	∀-9
	Base Location	Difference	Base Location	Difference
	:+			77.41

	Base Locati n	Difference	Base L cation	Difference
	114250	A DEL	176222	T-C
	115217	C-G	176524	A-T
	117995	G-A	176684	G-A
_	118874	A-G	176815	T-C
5	119470	T-C	177049	T-C
	119646	G-T	177065	G-T
	120853	C-T	178285	T-C
	121582	G-A	178551-178552	CTITITITITITINS
	123576	A-C	179114-179115	AINS
10	125581	C-T	179260	C-G
	125970	G-T	179281	C-G
	126197	A-G	180023	G-C
	126672	A DEL	180430	T-C
	126672	G-C	180773	T-C
15	128220-128221	AINS	180824	Ť-C
	132569	C-T	181097	C-T
	133572	A-C	181183	A-T
	134064	T-G	182351	C-T
	136999	G-A	183197	G-A
20	137784	C-T	183623	A-T
	138903	G-A	183653	G-T
	139159-139160	AINS	183657	T-G
	140359	G-A	183795-183796	AINS
	140898	IC-T	184060	G-A
25	141313	CDEL	184993	G-A
	141343	T-C	185918	A-G
	142148	T-C	186036	T-C
	142178	C-A	186506-186507	
	142433-142434	ATAGA INS	186561-186568	TAAC INS
30	143783	C-T	186690	TATTTATT DEL
	144090	С-Т	186751	G DEL
	144220-144221	AINS	187221	T-A
	144725	A-C	187260	A-G
	145732-145733	AAAAAAAAAAAAA INS	187444-187447	A-G
35	147016-147017	CG DEL	187831-187832	CTCT DEL
	147021	IG-T	188638	CINS
	147536	T-G		G-A
	148936	T-A	188642	C-T
	149061	T-C	189246	T-C
40	154341	A-T	190340	A-C
	154588	G-A	190354	A-G
	155464	IG-A	190762	A-G
	158574	C-G	191260	G-T
	160007	C-T	193018-193019	AGAT INS
45	164348		193147	T-G
45	164499	A-T	193196-193197	CINS
	166677-166678	C-G	193499	C-T
	167389	AAAG INS	193738	C-G
	168506-168507	G-A	193984-193985	ACACACAC INS
50	168515	AGGATGGTCT INS	194064	C-G
30		T-C	194504	A DEL
	169413-169414	AA INS	194734	G-A
	170300-170301 170491	TTGTTGTTGTTG INS	194890	A-C
		G-A	195404	G-A
55	173428	T-C	195693	A-T ,
JJ	173642	G-A	196205	G-A
	173948	T-G	197424	C-T
	175330 175836	T-C	197513	C-T
	176200	T-C	197670	G-A
	110200	G-C	198055	C-A

SUBSTITUTE SHEET (RULE 26)

Frequency of unaffected varian	Frequency of ancestral variant in	L cation	
esm comordo mobrisi ni	rand m chromosomes]
% 4 7	%ES	219560	
%SE	%99 %99	776412	
%0S	%0S	202112	
%2V %9L	%t5	2145495	S
%Lt	%\$9 %£\$	201712	_
%2 <i>\</i>	%ES	210299	
50%	%08		
25%	%8¢	208634	
%SZ	%SZ	207400	0
%09	%09	205284	1
%47	%ES	204341	1
%ZV	%8S	202880	l
%Z	%95 %96	200027 202662	9
%Z† %SL	%8S	120003	_
%SÞ	%SS	198692	
%Sb	%SS	104891	
%St	%\$\$	SS0861	
%0 7	%09	195633	0
%\$4	%SZ	195404	
%57	%99 ***********************************	068461	1
%17	%28 %ES	175330	1
%L1	%55 %55	849571	, ع
%0Z \$2%	%08 %SS	173642	S
%0Z	<u>%08</u>	515891	Į
%78	%81	700091	ļ
%ZÞ	%85	19061	
%8i	%Z8	148936	0
%0	%00l	147536]
% * 5	%9 <i>†</i>	147021	1
%SÞ	%\$\$	141343	!
%5V %5V	%95 %95	13803) S
%6l %St	%18 %SS	138903	S
%28	%81	125581	ł
50%	%08	121582	l
82%	%81	120853	1
%SI	%58	478811	0
%09	%0S	115217] .
%09	%0¢	113130	1
%ZS	%87 %87	130011	1
%05 %Z\$	%05 %87	828701	Si
%S7 %O2	72% 20%	747£01 31£36	
50%	%08 %C7	46116	{
%9Z	%SZ	88006	
%09	%09	82798	
%0S	%0S	SÞ968	0
%LE	%E9	88528]
%97	%SZ	26878]
%OÞ	%09	£1778	
%0S	%05 %09	22978	3:
20% 51%	%0S - %64	\$8698 50738	S
		60.460	

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2:159; 68-1:167; 241-5:108; 241-29:113; 373-8:151; and 373-29:113, D6S258:199, D6S265:122, D6S105:124; D6S306:238; D6S464:206; and D6S1001:180.

Table 2 lists the frequency of about 100 of the alleles defined by the polymorphic sites of the invention in the general population. As is evident from the Table, certain of these alleles are present rarely in the general population. These polymorphisms are thus preferred as surrogate markers in diagnostic assays for the presence of a mutant HFE allele ("gene mutation") such as 24d1 or 24d2. Preferably, the frequency of the polymorphic allele used in the diagnostic assay in the general population is less than about 50%, more preferably less than about 25%, and most preferably less than about 5%. Thus, of the genotypes defined by the alleles listed in Table 2, polymorphisms occurring at base 35983 and base 61465 of Figure 1 are preferred.

It will be understood by those of skill in the art that because they were identified in an ancestral HH homozygote, the haplotypes defined by the polymorphic sites of Table 1 are predictive of the likely presence of the HFE gene mutation 24d1. Thus, for example, the likelihood of any affected individual having at least two or more of any of the polymorphic alleles defined by Table 1 is greater than that for any unaffected individual. Similarly, the likelihood of any affected individual having at least three or more of any of the polymorphic alleles defined by Table 1 is greater than that for any unaffected individual.

Thus, for example, in a diagnostic assay for the likely presence of the HFE gene mutation in the genome of the individual, DNA or RNA from the individual is assessed for the presence or absence of a haplotype of Table 1, wherein, as a result, the absence of a haplotype of Table 1 indicates the likely absence of the HFE gene mutation in the genome of the individual and the presence of the haplotype indicates the likely presence of the HFE gene mutation in the genome of the individual.

The markers defined by the polymorphic sites of Table 1 are additionally useful as markers for genetic analysis of the inheritance of certain HFE alleles and other genes which occur within the chromosomal region corresponding to the sequence of Figure 9 which include, for example, those disclosed in copending U.S.S.N. 08/724,394.

As the entire nucleotide sequence of the region is provided in Figure 9, it will be evident to those of ordinary skill in the art which sequences to use as primers or probes for detecting each polymorphism of interest. Thus, in some embodiments of the invention, the nucleotide sequences of the invention include at least one oligonucleotide pair selected from the sequence of Figure 9 or its complement for amplification of a polymorphic site of Table 1. Furthermore, in some embodiments of the invention a preferred hybridization probe is an oligonucleotide comprising at least 8 to about 100 consecutive bases from the sequence of Figure 9, or the complement of the sequence, wherein the at least 8 to about 100 consecutive bases includes at least one polymorphic site of Table 1. In some embodiments the polymorphic site is at base 35983 or base 61465.

It will also be appreciated that the nucleic acid sequences of the invention include isolated nucleic acid molecules comprising about 100 consecutive bases to about 235 kb substantially identical to the sequence of Figure 9, wherein the DNA molecule comprises at least one polymorphic

5:108; 241-29:113; 373-8:151; and 373-29:113, alleles D6S258:199, D6S265:122, D6S105:124, D6S306:238, D6S464:206; and D6S1001:180, and/or alleles associates with the HHP-1, the HHP-19 or HHP-29 single base-pair polymorphisms can also be used to assist in the identification of an individual whose genome contains 24d1 and/or 24d2. For example, the assessing step can be performed by a process which comprises subjecting the DNA or RNA to amplification using oligonucleotide primers flanking at least one of the base-pair polymorphisms HHP-19, 24d2, oligonucleotide primers flanking at least one of the base-pair polymorphisms HHP-1, HHP-19, and HHP-29, oligonucleotide primers flanking at least one of the microsatellite repeat alleles, or oligonucleotide primers for any combination of polymorphisms or microsatellite repeat alleles, or Oligonucleotides useful in diagnostic assays are typically at least 8 consecutive nucleotides in length, and may range upwards of 18 nucleotides in length, and may range upwards of 18 nucleotides in length, and may range upwards of 18 nucleotides in length, and may range upwards of 18 nucleotides in length, and may range upwards of 18 nucleotides in length, and may range upwards of 18 nucleotides in length, and may range upwards of 18 nucleotides in length, and may range upwards of 18 nucleotides in length, and may range upwards of 18 nucleotides in length.

nucleotides in length, and may range upwards of 18 nucleotides in length to greater than 100 or more consecutive nucleotides. Such oligonucleotides can be derived from either the genomic DNA of Figure 8 or 9, or cDNA sequences derived therefrom, or may be synthesized.

Additionally, the proteins encoded by such cDNAs are useful in the generation of antibodies for analysis of gene expression and in diagnostic assays, and in the purification of related proteins.

E. General Methods

The nucleic acid compositions of this invention, whether RNA, cDNA, genomic DNA, or a hybrid of the various combinations, may be isolated from natural sources, including cloned DNA, or may be synthesized in vitro. The nucleic acids claimed may be present in transformed or transfected whole cells, in a transformed or transfected cell lysate, or in a partially purified or substantially pure form.

Techniques for nucleic acid manipulation of the

nucleic acid sequences of the invention such as subcloning nucleic acid sequences encoding polypeptides into expression vectors, labeling probes, DNA hybridization, and the like are described generally in Sambrook et al., Molecular Cloning - a Laboratory Manual (2nd Ed.), Vol. 1-3, Cold Spring Harbor, New York, (1989), which is incorporated herein by reference. This manual is hereinafter referred to as "Sambrook et al."

There are various methods of isolating the nucleic acid sequences of the invention. For example, DNA is isolated from a genomic or cDNA library using labeled oligonucleotide probes having sequences complementary to the sequences disclosed herein. Such probes can be used directly in hybridization assays. Alternatively probes can be designed for use in amplification techniques such as PCR.

To prepare a cDMA library, mRMA is isolated from tissue such as heart or pancreas, preferably a tissue wherein expression of the gene or gene family is likely to occur. cDMA is prepared from the mRMA and ligated into a recombinant vector. The vector is transfected into a recombinant host for propagation, screening and cloning. Methods for making and screening cDMA libraries are well known. See Gubler, U. and Hoffman, B.J. Gene 25:263-269 (1983) and Sambrook et al.

F ragenomic library, for example, the DMA is extracted from tissue and either

mechanically sheared or enzymatically digested to yield fragments of about 12-20 kb. The fragments

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high level expression of a cloned gene, it is desirable to construct expression plasmids which contain, at the minimum, a strong promoter to direct transcription, a ribosome binding site for translational initiation, and a transcription/translation terminator. The expr ssion vectors may also comprise generic expression cassettes containing at least one independent terminator sequence, sequences permitting replication of the plasmid in both eukaryotes and prokaryotes, i.e., shuttle vectors, and selection markers for both prokaryotic and eukaryotic systems. See Sambrook et al. Examples of expr ssion of ATP-sensitive potassium channel proteins in both prokaryotic and eukaryotic systems are described below.

a. <u>Expression in Prokaryotes</u>

A variety of procaryotic expression systems may be used to express the proteins of the

al. for details concerning selection markers for use in E. coli.

invention. Examples include E. coli, Bacillus, Streptomyces, and the like.

It is preferred to construct expression plasmids which contain, at the minimum, a strong promoter to direct transcription, a ribosome binding site for translational initiation, and a transcription/translation terminator. Examples of regulatory regions suitable for this purpose in *E. coli* are the promoter and operator region of the *E. coli* tryptophan biosynthetic pathway as described by Yanofsky, C., J. Bacteriol. 158:1018-1024 (1984) and the leftward promoter of phage lambda (Pλ) as described by Herskowitz, I. and Hagen, D., Ann. Rev. Genet. 14:399-445 (1980). The inclusion of selection markers in DNA vectors transformed in *E. coli* is also useful. Examples of such mark its include genes specifying resistance to ampicillin, tetracycline, or chloramphenicol. See Sambrook et

To enhance proper folding of the expressed recombinant protein, during purification from *E. coli*, the expressed protein may first be denatured and then renatured. This can be accomplished by solubilizing the bacterially produced proteins in a chaotropic agent such as guanidine HCI and reducing all the cysteine residues with a reducing agent such as beta-mercaptoethanol. The protein is then renatured, either by slow dialysis or by gel filtration. *See* U.S. Patent No. 4,511,503.

Detection of the expressed antigen is achieved by methods known in the art as radioimmunoassay, or Western blotting techniques or immunoprecipitation. Purification from *E. coli* can be achieved following procedures such as those described in U.S. Patent No. 4,511,503.

b. Expression in Eukaryotes

A variety of eukaryotic expression systems such as yeast, insect cell lines, bird, fish, and mammalian cells, are known to those of skill in the art. As explained briefly below, a sequence of interest may be expressed in these eukaryotic systems.

Synthesis of heterologous proteins in yeast is well known. <u>Methods in Yeast Genetics</u>, Sherman, F., et al., Cold Spring Harbor Laboratory, (1982) is a well recognized work describing the various methods available to produce the protein in yeast.

Suitable vectors usually have expression control sequences, such as promoters, including 3-phosphoglycerate kinase or other glycolytic enzymes, and an origin of replication, termination sequences and the like as desired. For instance, suitable vectors are described in the literature (Botstein, et al., Gene 8:17-24 (1979); Broach, et al., Gene 8:121-133 (1979)).

Saveria-Campo, M., 1985, "Bovine Papilloma virus DNA a Eukaryotic Cloning Vector" in <u>DNA Cloning</u> Vol. II a Practical Approach Ed. D.M. Glover, IRL Press, Arlington, Virginia pp. 213-238.

The host cells are competent or rendered competent for transformation by various

means. There are several well-known methods of introducing DNA into animal cells. These include: DNA, treatment of the recipient cells with liposomes containing the DNA, DEAE dextran, electroporation and micro-injection of the DNA directly into the cells.

The transformed cells are cultured by means well known in the art (Biochemical Methods in Cell Culture and Virology, Kuchler, R.J., Dowden, Hutchinson and Ross, Inc., (1977)). The expressed polypeptides are isolated from cells grown as suspensions or as monolayers. The latter are recovered by well known mechanical, chemical or enzymatic means.

2. Purification

The proteins produced by recombinant DNA technology may be purified by standard techniques well known to those of skill in the art. Recombinantly produced proteins can be directly expressed or expressed as a fusion protein. The protein is then purified by a combination of cell lysis (e.g., sonication) and affinity chromatography. For fusion products, subsequent digestion of the fusion protein with an appropriate proteolytic enzyme releases the desired polypeptide.

The polypeptides of this invention may be purified to substantial outsity by a continuous protein.

The polypeptides of this invention may be purified to substantial purity by standard techniques well known in the art, including selective precipitation with such substances as ammonium sulfate, column chromatography, immunopurification methods, and others. See, for instance, Roces, Protein Purification: Principles and Practice, Springer-Verlag: New York (1982), incorporated herein by reference. For example, in an embodiment, antibodies may be raised to the proteins of the invention as described herein. Cell membranes are isolated from a cell line expressing the recombinant protein, the protein is extracted from the membranes and immunoprecipitated. The recombinant protein, the protein is extracted from the membranes and immunoprecipitated. The proteins may then be further purified by standard protein chemistry techniques as described above.

3. Antibodies

As mentioned above, antibodies can also be used for the screening of polypeptide products encoded by the polymorphic nucleic acids of the invention. Such antibodies can be utilized in a variety of other contexts in accordance with the present invention. Such antibodies can be utilized for the diagnosis of HH and, in certain applications, targeting of affected tissues.

Thus, in accordance with another aspect of the present invention a kit is provided that

is suitable for use in screening and assaying for the presence of polypeptide products encoded by the polymorphic nucleic acids of the invention by an immunoassay through use of an antibody which specifically binds to polypeptide products encoded by the polymorphic nucleic acids of the invention in combination with a reagent for detecting the binding of the antibody to the gene product.

Once hybridoms cell lines are prepared, monoclonal antibodies can be made through

conventional techniques of priming mice with pristane and interpentioneally injecting such mice with the hybrid cells to enable harvesting of the monoclonal antibodies from ascites fluid.

In connection with synthetic and semi-synthetic antibodies, such terms are intended to cover antibody fragments, isotype switched antibodies, humanized antibodies (mouse-human, human-cover antibody fragments, isotype switched antibodies, humanized antibodies (mouse-human, human-cover antibody fragments, isotype switched antibodies, humanized antibodies (mouse-human, humanized antibody fragments).

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The number of clones per DS contig varied between 1 to 22 with the length of each contig ranging from 250bp to 850 bp. Small sequence-tag-sites PCR assays wire developed for each DS contig and two experiments were carried out concomitantly; mapping each DS contig back to the bacterial clon contig of the region and testing for the presence of each DS contig in cDNA libraries. Overall, 86 or 80% of the DS contigs mapped back to the region and were found to be in cDNA libraries. The number of 80% mapping to the region was probably an underestimate of the fidelity of the direct-selection sinc PCR assays which cross exon-intron boundaries would be expected to fail or give larger size products, thereby being scored negative.

b. Exon-Trapping

CsCI-purified genomic P1 (Genome Systems), BAC (Research Genetics) and PAC (Genome Systems) DNAs were digested with BamHI, Bgl II, Pst I Sac 1 and Xho I and 125 no of each digest ligated into 500 ng pSPL3 (Church et al. Nature Genetics 6:98-105 (1994)) (Life Technologies, Gaithersburg, MD) digested with the appropriate restriction enzyme and phosphatased with calf intestinal alkaline phosphatase (USB, Cleveland, OH). One tenth of the ligation was used to transform XL1-Blue MRF' cells (Stratagene, La Jolla, CA) by electroporation. Nine tenths of the electroporation was used to inoculate 10 ml of LB + 100µg/ml of carbenicillen and after overnight growth, DNA was prepared using Qiagen Q-20 tips (Qiagen GmbH, Hilden Germany). The remaining one tenth was plated on LB +100 µg/ml carbenicillen plates to evaluated the efficiency on cloning and to test individual clones for the present of single inserts. COS-7 cells were seed overnight at a density of 1.4 x10⁵/well in 6 well dishes. One ug of DNA was transfected using 6ml of Lipofect-Ace. Cytoplasmic RNA was isolated 48 hr post-transfection. RT-PCR was carried out as described by Church et al. (ibid) using commercially available reagents Life Technologies, Gaithersburg, MD). The resulting CUA-tailed PCR fragments for each restriction digested bacterial clone were pooled and UDG cloned into pSP72-U (a derivative of pSP72). The DNA was transformed in DH5α and the cells plated onto nylon membranes. After overnight growth, duplicates were made and the DNA hybridized to ³²P end-labeled oligos designed to detect various background products associated with the pSPL3 vector. One set of filters was hybridized with the following gel-purified oligos in 6X SSC aqueous hybridization solution at 42° C:

vector-vector splicing

5'-CGACCCAGCAACCTGGAGAT-3'

cryptic donor-1021

5'-AGCTCGAGCGGCCGCTGCAG-3'

cryptic donor-1134

5'-AGACCCCAACCCACAAGAAG-3'

The filters were washed twice in 6X SSC, 10 mM sodium pyrophosphate (NaPPi) at 60°C, 30 mins.

After overnight autoradiography, non-hybridizing clones were picked and grown in 250 µl of LB + 100µg/ml of carbenicillin in 96 well mini-rack tubes. The samples were analyzed by PCR using the secondary PCR primers supplied in the kit (Life Technologies, Gaithersburg, MD) and those clones with inserts greater than 200 bp were selected for sequencing.

Ninety-six exon traps per bacterial clone were sequenced for a total of 768 reactions and the resulting data analyzed by BLAST. In addition, each potential exon was searched against a database of the 86 DS contigs to eliminate redundant sequences. PCR assays were diveloped for

yi77b02 b132a12 NSH Alue - CDNA 37 yh76b03 b132a12 NSH Alue -	35
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yh76b05 b132a12 NSH Alue -	
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\\ \text{A638C0} \\ \text{A638C0} \\ \text{P50P20}	
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Vh76b05 b132a12 NSH Alu	
VI V DOZ DIBZEIZ NCH NCH CDNV 37	
ym29g03 b132a12 Histone H2A NSH + cDNA 37	12
+ - UIA HZN SIBSEID IIDTEDY	1
y804f09 b132b12 Line element Alu - +	J
HUM160hiib bi32ai2 none na na na	
H2CSEE087 P137915 us M2H +	
ybl2h11 b132a12 NSH Histone H3.1	10
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yh54f11 p196e20 none NSH	
AC18E10 PC45P21 NSH na na	
ynOlcO5 p196£20 BUTYBOVIN na na cDNA 32	
7504h08 pc45p21 nsh + - cDnA 44	S
ye25803 p196e20 BUTYBOVIN NSH na cDNA 44	5
yd17d06 p196e20 NSH na na cDNA 23	
yv88c09 p196e20 BUTYBOVIN na na cDNA 29	
67 ANG and	
Clone name Bacterial Homology 5' Poly A+ Genomic cDNA clone Diastx blastx signal' poly (A) _{o4} Homology yo65f06	

d. cDNA library screening

Technologies, Gaithersburg, MD. Colonies were plated on Hybond N filters (Amersham) using Superscript plasmid cDNA libraries, brain, liver and testis, were purchased from Life

cloned were not used in any screen. Therefore, it is possible that some additional genes within this 1 megabase region may have escaped detection.

A list of thes cDNAs cloned and a comparison of the methods used to find them is presented in Table 4. Direct selection found 14 out of the 18 cDNAs contained within the boundaries of the YAC used in the experiment. Exon trapping found 15 out of the 19 cDNAs contained within the boundaries of the large insert bacterial clone contig. Sample sequencing identified 11 genes that had corresponding ESTs in the public database.

Table 4. Comparison of gene finding methods

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157c 28 zinc finger EST03556 2 1 157c3 30 nonhistone yv81d05 1 none 157c3 46 ORF yd88g11 1 157c3 20 BT none none 3 p18696 21 BTF1 yn01G5 4 5 yg23d08 yg57h09 yu15h03 4 45p21 32 BTF2 yg78f10 7 3 yn01c05 45p21 29 BTF3 ye25g03 2 9	
157c3 30 nonhistone yv81d05 1 none yvh07a10 157c3 46 ORF yd88g11 1 15 157c3 20 BT none none 3 p18696 21 BTF1 yn01G5 4 5 yg23d08 yg57h09 yu15h03 45p21 32 BTF2 yg78f10 7 3 yn01c05	on Trap
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yu15h03 45p21 32 BTF2 yg78f10 7 3 yn01c05	
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yo65f06	
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20 45p21 44 BTF5 ys04h08 2 4	
3e17 41 genomic? none none 1	
132a2 43 genomic? none none 3	
132a2 36 genomic? none 1 none	
132a2 37 histone 2A ym29g03 3 none	
yh87a03	•
25 75114 24 MHC class 1 ye98g01 1 2	
132a2 39 genomic? none none 4	
132a2 27 Ro/SSA none 3 4	
132a2 22B NPT1-like yr42a05 1 7	-
yf09g06	
20h20 22E NPT1-like none 2 5	
30 20h20 NPT1 NPT1 yp74c05 N/A 3	

display varying degrees of homology to BT. BTF1 (cDNA 21), BTF2 (cDNA 32), BTF5 (cDNA 44), and BTF3 (cDNA 29) are 45%, 48%, 46%, and 49%, identical to BT, whereas BTF4 (cDNA 23), which is more similar to BTF3 (cDNA 29), is only 26% identical. This low degree of identity to BT is largely due to a truncation at the carboxyl terminus of the protein. The BTF family falls into two groups: BTF1 and 2 which are more related to each other than to BT or the other BTF members, and BTF5, 3 and 4, which appears to have a common evolutionary origin. The order of these genes on the chromosome suggests that the BT gene has duplicated two times, giving rise to BTF1 and BTF5. Subsequently, it appears likely these two genes experienced further duplication events to give rise to the other appears in their groups.

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The three major components of BT, the B-G immunoglobulin superfamily domain (containing the V consensus sequence) (Miller et al. Proc. Natl. Acad. Sci. U.S.A. 88:4377-4381 (1991)), the transmembrane region, and the B30-2 exon are found in all of these proteins (with the exception of BTF4 (cDNA 23) which lacks the B30-2 exon by virtue of the carboxyl terminal truncation). The exon B30-2 is a previously noted feature of the MHC class 1 region found approximately 200 kb centrometic to the HLA-A gene (Vernet et al., J. Mol. Evol. 37:600-612 (1993)). In addition this exon is found in several genes of diverse function telomenc to HLA-A namely MOG (approximately 200 kb) and RFP (approximately 1 megabase) (Amadou et al. Genomics 26:9-20 (1995)).

The expression of the BTF genes fell into two patterns. BTF1 and BTF2 were expressed as a single major transcript of 2.9 kb and one minor transcript of 5.0 kb. These genes were expressed at high levels in all the tissues tested with the exception of the kidney where the expression level was less. The two genes are 90% identical at the DNA sequence level, therefore, it is possible that the signal observed on the northerns was the result of cross-hybridization and only one of the two genes was actually expressed. To address this possible tissue dependent expression that would suggest that both different tissues in order to detect possible tissue dependent expression that would suggest that both genes are expressed. Identical, and thus equivocal, results were obtained with both BTF1 and BTF2 genes are expressed. Identical, and thus equivocal, results were obtained with both BTF1 and BTF2.

The second group of genes, BTF3-5, are expressed as three (BTF5) (Figure 4A) and the second group of genes, BTF5 is expressed at moderate levels in all bissues tested with the exception of the kidney where the expression level is less. RT-PCR experiments showed that mRNA from the BTF5 gene can be found in all tissues tested, including the kidney (Figure 4B). Identical results were obtained with primers from the other genes of this group (data not shown). These genes are also 90% identical to each other at the DNA sequence level (but only 58% identical to BTF1 and 2), hence like BTF1 and BTF2, cross-hybridization could account for the similarity in size and patterns on the northern blots and RT-PCR. This might be particularly true for the similarity in size and patterns on the northern blots and RT-PCR. This might be particularly true for the similarity in size and patterns on the northern blots and RT-PCR. This might be particularly true for the similarity in size and patterns on the northern blots and RT-PCR. This might be particularly true for the similarity in size and patterns on the northern blots and RT-PCR. This might be particularly true for the similarity in size and patterns on the northern blots and RT-PCR. This might be particularly true for the similarity in size and patterns on the northern blots and RT-PCR. This might be particularly true for the similarity in size and patterns on the northern blots are the size that a size and patterns on the northern blots are size and size than a size and patterns on the northern blots are size and patterns on the northern blots and RT-PCR.

ii. A gene with similarity to 52 kD Ro/SSA auto-antigen

58% amino acid similarity to the 52 kD Ro/SSA protein, an auto-antigen of unknown function that is frequently recognized by antibodies in patients with systemic tupus and Sjogren's syndrome (Anderson

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subset of the polymorphic alleles so defined were further studied to determine their frequency in a collection of random individuals.

The cell line HC14 was deposited with the ATCC om June 25, 1997, and is designated ATCC CRL-12371.

a. Cosmid Library Screening

The strategy and methodology for sequencing the genomic DNA for the affected individual was essentially as described in copending U.S.S.N. 08/724,394, hereby incorporated by reference in its entirety. Basically, a cosmid library was constructed using high molecular weight DNA from HC14 cells. The library was constructed in the supercos vector (Stratagene, La Jolla, CA). Colonies were replicated onto Biotrans nylon filters (ICN) using standard techniques. Probes from genomic subclones used in the generation of the sequence of the unaffected sequence disclosed in 08/724,394 were isolated by gel electrophoresis and electroporation. Subclones were chosen at a spacing of approximately 20 kb throughout the 235 kb region. The DNA was labeled by incorporation of 32P dCTP by the random primer labeling approach. Positively hybridizing clones were isolated to purity by a secondary screening step. Cosmid insert ends were sequenced to determine whether full coverage had been obtained, and which clones formed a minimal path of cosmids through the 235 kb region.

b. Sample Sequencing

A minimal set of cosmid clones chosen to cover the 235 kb region were prepped with the Qiagen Maxi-Prep system. Ten micrograms of DNA from each cosmid preparation were sonicated in a Heat Systems Sonicator XL and end-repaired with Klenow (USB) and T4 DNA polymerase (USB). The sheared fragments were size selected between three to four kilobases on a 0.7% agarose gel and then ligated to BstXI linkers (Invitrogen). The ligations were gel purified on a 0.7% agarose gel and cloned into a pSP72 derivative plasmid vector. The resulting plasmids were transformed into electrocompetent DH5α cells and plated on LB-carbenicillin plates. A sufficient number of colonies was picked to achieve 15-fold clone coverage. The appropriate number of colonies was calculated by the following equation to generate a single-fold sequence coverage: Number of colonies = size of bacterial clone (in kb)/average sequence read length (0.4 kb). These colonies were prepped in the 96-well Qiagen REAL, and the 5' to 3' DNA Prep Kit, and AGCT end-sequenced with oligo MAP1 using standard ABI Dye Terminator protocols. MAP1 was CGTTAGAACGCGGCTACAAT.

c. Genomic Sequencina

The MAP1 sequences from the cosmid clones HC182, HC187, HC189, HC195, HC199, HC200, HC201, HC206, HC207, and HC212 were assembled into contigs with the Staden package (available from Roger Staden, MRC). A minimal set of 3 kb clones was selected for sequencing with oligo labeled MAP2 that sits on the opposite end of the plasmid vector. The sequence of MAP2 was GCCGATTCATTAATGCAGGT. The MAP2 sequences were entered into the Staden database in conjunction with the MAP1 sequences to generate a tiling path of 3 kb clones across the region. The plasmid 3 kb libraries were concurrently transformed in 96 well format into pox38UR (available from C. Martin, Lawrence Berkeley Laboratories). The transformants were subsequently mated with JGM (Strathman et al. P.N.A.S. 88:1247-1250 (1991) in 96 well format. All matings of the

PCR primers for detection:

182.1G7.F 5'-GCATCAGCGATTAACTTCTAC -3' 182.1G7.R 5'-TTGCATTGTGGTGAAATCAGGG -3'

For the detection assay, the biotinylated primers used were as follows.

182.1G7.C 5' (b)CTGAGTAATTGTTTAAGGTGC-3'

182.1G7.T 5' (b)CTGAGTAATTGTTTAAGGTGT-3'

The phosphorylated digoxigenin-labeled primer used was:

187.151. St. (p)AGAGAGATAGATAGATAGGG -3"

A further rare single base pair change was detected at 61,465bp. The inheritance pattern of this polymorphism, C195.1H5C/T (a G to A change on the opposite strand), is identical to that of 24d1. The frequency of T occurring at that position (C195.1H5T) observed in a set of 76 patients was 78.5% as compared to 5% in random individuals.

15 PCR primers for detection:

1951H5.3F 5'-CAACTGAATATGCAGAAAAAGTACACC-3' 8'-CAACTGAATATGCAGAAAAAAAGTACACC-3'

For the detection assay, the biotinylated primers used were:

1951H5.3.4 5' (b)AGTAGCTGGGACTCACGGTGT-3'

1957H5.3.5 5' (b)AGTAGCTGGGACTCACGGTGC-3'

The phosphorylated digoxigenin-labeled primer used was:

1951H5.3.6 5' (p)GCGCCCCCCCCCCCCCCCCT-3'

These rare alleles are thus preferred surrogate markers for 24d1 and are especially

25 useful in screening assays for the likely presence of 24d1 and/or 24d2.

All publications, patents, and patent applications cited herein are hereby incorporated

by reference in their entirety.

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6	1:206, 65-2:159, 68-1:167, 241-5:108, 241-29:113, 373-8:151, 373-29:113, D6S258:199, D6S265:1
7	D6S105:124, D6S306:238, D6S464:206, or D6S1001:180.
1	11. The method of claim 9, wherein the haplotype comprises at least two polymorphic
2	sites of Table 1.
1	12. The method of claim 11, wherein one of the at least two polymorphic sites of Table
2	is at base 35983 or 61465.
1	13. The method of claim 11, wherein the haplotype comprises at least three polymorph
2	sites of Table 1.
1	14. A method to determine the presence or absence of the common hereditary
2	hemochromatosis (HFE) gene mutation in an individual comprising:
3	providing DNA or RNA from the individual; and
4	assessing the DNA or RNA for the presence or absence of a genotype defined by a
5	polymorphic allele of Table 1,
6	wherein, as a result, the absence of a genotype defined by a polymorphic allele of Table 1
7	indicates the likely absence of the HFE gene mutation in the genome of the individual and the
8	presence of the genotype indicates the likely presence of the HFE gene mutation in the genome of the
9	individual.
1	15. The method of claim 15, wherein the polymorphic allele occurs in less than about 5
2	of a random population of individuals.
1	16. The method of claim 15, wherein the polymorphic allele occurs in less than about 2
2	of a random population of individuals.
1	17. The method of claim 15, wherein the polymorphic allele occurs in less than about 5
2	of a random population of individuals.
1	18. The method of claim 15, wherein the genotype is C182.1G7C or C195.1H5T.
1	19. A kit comprising one or more oligonucleotides of claim 1.
1	20. A kit comprising at least one oligonucleotide pair of claim 3.

A culture of lymphoblastoid cells having the designation ATCC CRL-12371.

21.

An antibody that is specifically immunor active with the polypeptide of claim 54.	.73	
A host cell stably transfected with the nucleic acid sequence of claim 52.	.95	
A vector comprising the nucleic acid sequence of claim 52.	.55	
The polypeptide encoded by the isolated nucleic acid sequence of claim 52.	* \$	
The isolated nucleic acid sequence of claim 52, wherein the nucleic acid is cDNA.	.53	
An isolated nucleic acid sequence comprising a sequence substantially identical to	b-3 25	TN :
An antibody that is specifically immunoreactive with the polypeptide of claim 48.	.re	
A host cell stably transfected with the nucleic acid sequence of claim 46.	·0S	1
A vector comprising the nucleic acid sequence of claim 46.	·6 †	1
The polypeptide encoded by the isotated nucleic acid sequence of claim 46.	.84	1
The isolated nucleic acid sequence of claim 46, wherein the nucleic acid is cDNA.	.74	ı
An isolated nucleic acid sequence comprising a sequence substantially identical to	.E2 46	t 18 S
An antibody that is specifically immunoreactive with the polypeptide of claim 42.	·SÞ	ı
A host cell stably transfected with the nucleic acid sequence of claim 40.	.44.	ι
A vector comprising the nucleic acid sequence of claim 40.	43	ι
The polypeptide encoded by the isolated nucleic acid sequence of claim 40.	·ZÞ	
The isolated nucleic acid sequence of claim 40, wherein the nucleic acid is cDNA.	'l+	ι
	1 E4.	s 8.
An isolated nucleic acid sequence comprising a sequence substantially identical to	.O.P.	

i	, 75 .	An isolated nucleic acid sequence comprising at least 18 contiguous nucleotides
) -	substantially ide	ntical to 18 contiguous nucleotides of NPT3.

- 76. An isolated nucleic acid sequence comprising at least 18 contiguous nucleotides substantially identical to 18 contiguous nucleotides of NPT4.
- 1 77. An isolated nucleic acid sequence comprising at least 18 contiguous nucleotides substantially identical to 18 contiguous nucleotides of RoRet.

2/162

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BT BTF1 BTF2 BTF5 BTF3 BTF4	AGPPRRVGIFLDYESGDISFYNMNDGSDIYTFSNVTFSGPLRPFFCLWSSGKKPLTICPI KESLCRVGVFLDYEAGDVSFYNMRDRSHIYTCPRSAFSVPVRPFFRLGC-EDSPIFICPA KESLCRVGVFLDYEAGDVSFYNMRDRSHIYTCPRSAFTVPVRPFFRLGS-DDSPIFICPA PKPPKKVGVFLDYETGDISFYNAVDGSHIHTFLDVSFSEALYPVFRILTLEPTALSICPA PEPPRKVGIFLDYETGEISFYNATDGSHIYTFPHASFSEPLYPVFRILTLEPTALTICPI
BT BTF1 BTF2 BTF5	ADGPERVTVIANAQDLSKEIPLSPMGEESAPRDADTLHSKLIPTQPSQGAP LTGANGVTVPEEGLTLHRVGTHQSL LTGASGVMVPEEGLKLHRVGTHQSL
BTF3 BTF4	PKEVESSPDPDLVPDHSLETPLTPGLANESGEPQAEVTSLLLPAHPGAEVSPSATTNQNH
вт	
BTF1	
BTF2	
BTF5	
BTF3	KLQARTEALY
BTF4	

Figure 3 (Page 2 of 2)

SUBSTITUTE SHEET (RULE 26)

B30-2 DOMAIN	kD Ro Ret	52 kD RoRet
SALELLQEVIIVLERSESWNLKDLDITSPELRSVCHVPGLKKMLRTCAVHITLDPDTANPWLILSEDRRQVRLGDTQ SAQKLLQNVNDTLSRSWAVKLETSEAVSLELHTMCNVSKLYFDVKKMLRSHQVSVTLDPDTAHHEL!LSEDRRQVTRGYTQ ** *** * * * * * * * * * * * * * * * *	kD Ro	52 kD RoRet
© EVEIAIKRADWKKTVETQKSRIHAEFVQQKNFLVEEEQRQLQELEKDEREQLRILGEKEAKLAQQSQALQELISELDRRCHS KLSTAMRITKWKEKVQIQRQKIRSDFKNLQCFLHEEEKSYLWRLEKEEQQTLSRLRDYEAGLGLKSNELKSHILELEKKCQG	kD Ro	52 kD RoRet
O NIKKISQEAREGTQGERCAVHGERLHIFCEKDGKALCWVCAQSKKHRDHAMVPLEEAAQEYQEKIQVALGEIRRKQEIAEKI ALKKTDQEMSCEEHGEQFHIFCEDEGQLICWRCERAPQHKGHTTALVEDVCQGYKEKIQKAVTKIKQIEDRCTEQ *** ** ** ** ** *** *****************	kD Ro	52 kD RoRet
O MASAARLIMMWEEVTCPICLDPFVEPVSIECGHSFCQECISQVGKGGGSVCPVCRQRFLLKNLRPNRQLAMMVN MASTTSTKKMMEEATCSICLSLMTNPVSINCGHSYCHLCITDFFKNPSQKQLRQETFCCPQCRAPFHMDSLRPNKQLGSLIE *** * * *** *** *********************	kD Ro Ret	52 kD RoRet

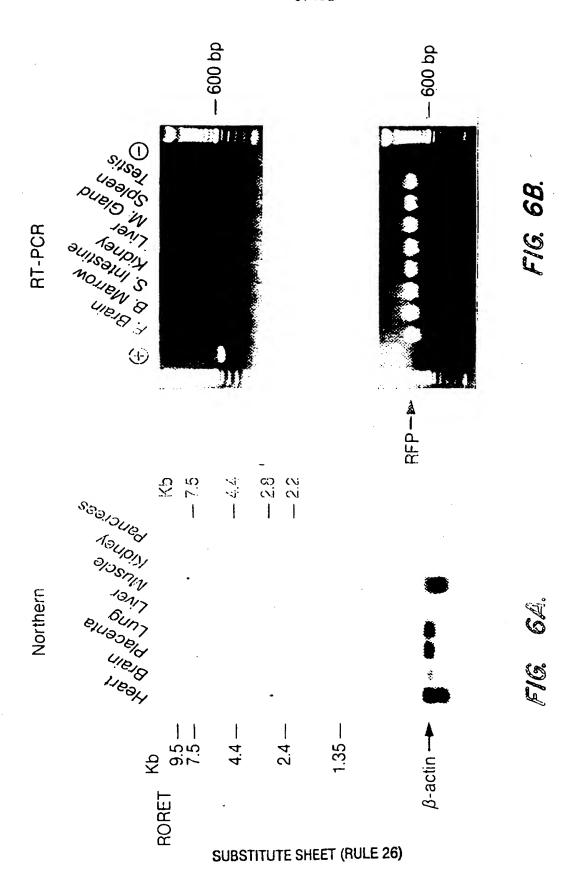
HLHEQPLLVGIFLDYEAGVVSFYNG-NTGCHIFTFPKASFSDTLRETTFQVYQYS----PLFLPPP--G----D--HLQVPPCQVGIFLDYEAGMVSFYNITDHGSLIYSFSECAFTGPLRPFFSPGFNDGGKNTAPLTLCPLNIGSQGSTDY

52 RoRet

ξD

Ro

-CYSTEINE-RICH DOMAIN



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Figure 7 (3 of 6)

Eigure 7 (5 of 6)

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481		, rwgrarcwal	TATIGAACTG	GGGAGAACYC	CARCCORMS.	
541	22	. WITCHIGIGI	CTTGGAAGTG	TTTACCCTC	77C7CCT70C	
601	GONGAIGI	CANGIGNAAA	TGTGGCTACA	CACATTCC		
661	* OOUGY I GIN	CHANTIGGAAG	TTTACTGCAT	TATACATACTC	TTTCCSSCC	
721	CHARGECATIA	AIGAGACAGA	ACAAAGACTA	GGGACCAGAG	CCDDCCCCD	>
781		INGININGIC	IGGICATTIT	GAGGTGA ATA	CTTAATAAAA	033000
841	TOMOTOTA	MATTIAGAGE	CCTACACTTT	TAGCTCTGAC	TATTAACCAA	TAGROGALA
901	MIGGNIAIG	GITHICIGGG	TGGTGTCTGT	GAAATAATT	ANGCCACCAN	G1 G1 magnes
961	CCAGAAACIG	ACTATGCTGG	CAACTTGGAT	$-CTT\DeltaC\DeltaTTTC$	CNCCCTCCNC	3.3.mm.a
1021	MAKE TANKET G.T.	CIAICGITTA	AGCCACCAGT	CTGTAGTATT	THETTATE	1000000
1081	OUCTWGIII	IGGIACCCAG	GCGTGGGATG	CTGCAACAAC	AAATACCTAA	A CA TCCCCA A
1141	0100011100	MANTIGGIGA	TGGGTAAAGG	CTGGAAGAGT	TTGAGGTTCA	TACTACABAA
1201	MOCCAMITGI	GAAGGGACTA	TTGAAAGAAA	TATGGACATT	AAAGGCAATT	CTCCCAAAGG
1261	CICAGAAAGG	AAGAGAGCTG	GACAGAAAGC	TTCCATTTTC	ATACAAACTT	
1321	CGATCATGGA	TAGAATATTA	AATATGCTGG	TTAAAATATG	GACTTTAGGC	CACCCCTCCT
1381	GGCTCACGCC	TGTAATCTCA	GCACTTTGGG	AGGCTGAGGG	CACAGATCAC	GACCTCCCCA
1441	GITTGAGACC	AGCCTGGCCA	ATATGGCGAA	ACCCTGTCTC	TACTADADAT	ACA A A A ATTA
1501	GCTGGGCATG	GTGATGTGCT	TCTGTGGTCC	CAGCTACTCG	GGAGGCTGAG	CCTCAACAATIA
1561	CGCTTAAACC	CGGGGGGTGG	AGGTTGCAGT	GACCCAAGAT	CACACCACTG	CACTCCACCC
1621	TGGGATACAG	AGCAGGACTC	CACTCCCCC	GCCACACACA	CACAAAAAA	TATICCAGCC
1681	GGACATTAAA	GTCAACTCTT	GTGAGGTCTC	AGATGAAAAT	GAGGGACAGG	ጥጥ እ ምጥር ርር እ እ እ
1741	CIGIAGAAAT	CACTGTTCTT	GTTACAATGT	GTCAAGAACT	TGGCTGAATT	ACCCTCTACT
1801	GITTACTGGA	AAGAACTTAT	AAGCAGTAAA	ACTGGATATT	TACCAGAAGA	GATCTCTAAC
1861	CAMAGIATIG	AAGGTGTGAT	TTAGGTCCTC	CTTACTGCTT	AAAGTGAAAT	GTGACACCAA
1921	AGAGCCGAAA	TAAAGAAGGA	ATTTTTAAGC	AAAACACAAT	CAGAACTTCC	AC ATTTCCCA
1981	TAGATTTCTC	AATCTATATT	GTAAAAATTG	AGAAAGTTTT	TCTTGAAGAG	CTATCCTTCA
2041	ACAATGTTTT	CITTITCTT	TTTTTTCTTG	GTTTTATTTT	TATTTTTATC	TTTTTTTT
2101	CAGGGTCTGG	CTATGTCATC	CAGGCTGGAG	TGCAGTGGCA	CAATCTCAGT	TCACTCCAAC
2161	CITIGCCTTC	AGGCTCAAGC	AATCCTCCCA	CCTCAGCCTC	CTAAGTAGCT	CCCACTACAT
2221	GTATGCACCA	CCACACCCTG	GCTAATTTTT	TGTTGTTGTT	TATAGAGATG	CCCTTTTCAC
2281	AIGITGCCTA	GGCTGGTCTC	TAACTCCTGA	GCTCAAGTGA	TCTGCCCTCC	TCACTCTCC
2341	AAAGTGTTGG	GATTACAGGC	GTGAAACACT	GAGCCTAGCC	TGAACAACCA	TTTCATABAG
2401	AGATAATGGG	TGTGACCCAA	GGATTTAATC	AGCCATCTCA	GCAGAAGCCA	GGAAGAGAGA
2461	IGGGATTATT	CCAGCAGAGA	CACTGCCAAT	TTAAACTAAC	GTAGGCAGAG	DADDCDCDAD
2521	GGAACAAAGG	AAGGTTGTCG	ACTTTTTGAA	TTCTATAGAA	CAGGATCATA	GAGCTACCTC
2581	GCTGTCAATG	TGTACTATTC	TTTAAGAAAA	GGAAAGACTG	ACCCACCAAA	GGCAACTTAC
2641	AAGATCACTA	GGGCTGACTC	TTTTGTTTTT	TCTTGAGGCA	GTCTCACTGT	CACCCAGGGT
2701	GIAGGGCAAT	GGTGTGATCT	CAGCTCACTG	CAATCTCCAC	CTCCCAGGTT	CAAGGGATTC
2761	ICTTGCCTTA	GACTCCCAAG	TAGCTGGGAT	TACAGGCTCT	AAATCTGTAC	CCTCCCGAGT
2821	AGCGCTCCTG	CCACCACTTG	CCCAGCTAAT	TTTTGTATTT	TTAGTAGAGA	TGGGGTTTCA
2881	CTATGTTGGC	CAGGCTAGTT	TGGAACTCCT	GACCTCCAGT	GATCCATTCT	CATTGGCCTC
2941	CCAAAGTGCT	GGGATTACAG	GCAGGAGCCG	CCAGGGCTGC	CACTTTGATG	TCAGACTCAG
3001	AGAGTACAGA	TGGGATAGGG	TGGGGGTGGG	AACATGTAGT	CAAGGCTGAC	TCTACCTGTT
3061	TCAAAGATGC	CCTGCAGAAC	TGTGTGGGAG	TCTCTCACAG	ATGGCTGCCT	GGGTGGGACC
3121	CCACCAAACT	GAAAGACCGA	GACTTCAGGC	AGGGCAGATG	GAGTAGGCCA	ACTACAGAGC
3181	CAGAGGTGAC	ACTGAGACAC	CACTGGGCCT	GGAAATCAGG	GCATCAAGCC .	AAAGAGGGTT

Figure 8 (Page 1 of 73)

SUBSTITUTE SHEET (RULE 26)

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13261	MOTOTINGGA	LIAACAGGAC	i ACAAAAAGGA	. AATCAAGGAA	ACCGAATCTC	TOOMS
13321	WACIGCIAIG	GCAGAGGCTC	: TACAGCTTAT	' ΤΑΥΓΑΔΤΉΤΤ	ACTA ATTMOS	01 mm1
13381	CCTTCACGTT	CITTAAGTAA	GGTTAGAGGA	CAGAAGAAAC	ATAATCTTCT	TACALAMORE
13441	VCINI IQAGI	CAGGGAAAA	AAAGAGTGCT	TTCAATATCT	GAATAAAACA	777777777
13501	VIIIICIWWA	CUTTAACGAG	TTTATTGTAA	GGGATGTGAT	GCTGGAAACT	ACCA A A CONA C
13561	AATTTTCTTC	TAAACTGAGA	ATCAGAATTA	TTCATATTCT	CAGCAGTGGT	CCCR CCTCR
13621	GGACTTCTGA	TCTTAATTAC	: ATACTTTTAT	TTCTTTAACT	GATCAACATC	CTABBERGE
13681	AACCTATGGC	TCTGTTTTTA	CCCACTTTAA	ATTCTGTTCT	ATTAGCACGG	TTACCTTCC
13741	IAATTGGCAA	TAAGATTGAG	ACTATCTTTT	TTTTTTTTTT	GAGACAGAAT	TTTCCTCTCT
13801	GGCCCAGGCT	GGGGTGCAGT	' GGCACAATCT	CGGCTCACTG	CAACCTCTGC	CTCCACCCTT
13861	CTAGCAATTT	TCCTGCCTCA	GCCTCCCCAG	TAGCTGGGAT	TACAGGTGCA	CCACCACCCC
13921	TGGCTAATTT	GTGCATTTTT	AGTAGAGATG	GGGTTTCGCC	ATGTTGGCCA	AACTGGTCTC
13981	GAACTCAGGT	GATCCACCTC	GGCCTCCCAA	AGTGATGAGA	TTACAGGCGT	GAGCCACCCT
14041	GCCCAGAAAA	GACTATCTTA	TTTTATGAAT	TTAAATAATT	GTGAAATTAT	CCACTTARGO
14101	GAATTAATAA	ATTATAATGT	AATCTTAAAT	TTTAGTTGGC	TTACATAAAG	ACTTANAGG
14161	CATCAATTTA	AATAAAAACT	CATTTGTCTA	AAAAAAATC	AAAAATTTTC	CTTGTGCTTT
14221	AAATGTGCTA	CCTCTTTAAG	TTCTAATTAA	GAGAAAAAA	GTTTAACTGT	GAGTTTCATT
14281	AGTGGTCTTA	GTTAACAGCT	TAAAGTATTT	TGTAAAAAAA	ATACTTCACA	הממדדדד
14341	AACTTAAAAA	TATTAATACC	TCTTTTATTA	GGTTTTTTTA	ATAAGGAAAA	TATATATAT
14401	ATCTAATCAA	GATTTTTTT	GGACAAATTG	GCTTAATAAT	TTCATTTTAA	AAATGGCTTC
14461	TTTATTCTTA	TACTGTAAAA	ATAATATTAG	CAGAATATTA	TAGTATACAC	AAGTTTAGGG
14521	TTCATATTCT	AAAAAACAAA	AACAAAAGCT	AATTTAACTT	GCATTTACTA	AATTTCTTCC
14581	ACTAGTTGTA	CTGGTTACAT	GAGTTAACAT	CACTTTATTT	ATTATTCTAA	AATTGTAAAT
14641	TATTCATTGA	ACCAAATTAA	ATGATAATAG	ATAATGTCAT	TTTTAAAATTT	GGAATTAAAT
14701	TTTATGTTAC	TAATTATAAG	GATTCAATGT	GTGAGCTTAA	GTACTGAGTT	CACAGTGTAT
14761	GATAACTTTA	AGAATTTAGG	TGAATATTAT	TAAATTGAGT	AAATTAATTC	TCAATCTTTC
14821	GATACCTGGA	CAATTTCTAA	ATTGGAGGGT	ACAAAATACA	AATCACAAGA	AACAGTGTAG
14881	TTTTATGCAA	ATAACATTTT	TACACAGTTT	AGAATAACCA	TTGATAAACA	GATAAGAGAA
14941	CATATGATTG	CCTTAGAATA	GATACTGTTG	CTTTCGCCAC	TTTAGATTTG	TAAATCACGT
15001	ACTGTATACG	TGTGGGCGTA	GAGGACCATG	CAGGTTTTGG	ATGACTGCCT	СТСТТТТССТ
15061	CATGCCTATG	CGGGAACACA	ATTGCCTGCT	TTGTTTAAGG	GCTATGGTTA	ATCCABACAG
15121	CTCTGACTCT	ATCAAGTACT	ATAGCTACAG	AGAAACACAA	GTAAGCATTC	GAGATAATGA
15181	CTACCTTGAG	CCTTTACTTA	TTTAAAAAGT	TGTTACTGTT	TGTTAATGTG	GTACATTCAA
15241	TTTACTATGG	ATTGTCACTC	TAAAATAAGA	CTTCAATCTT	TTTCTTATTT	TTATATAGCC
15301	ATGATTTATA	TTCATATCTT	AATGTAATAA	CCAATCTTCT	CTGACAACAT	TATAACAATG
15361	CTGGAACCTC	CATTTTCAGT	ACTTCAAACA	ACAAATACTG	CTTTTATACT	TCAGAGCAGA
15421	TGGATATGTG	CTTCCCAGTG	TAAACACATT	TGGAATCTCA	CTGAGAAATA	CACTATCACT
15481	AAAAATACAG	TTCTGAGATT	CATTAAAAGA	CCTCCAGAAT	TCTGGAAGTA	GGAAGTTTCC
15541	TCTTCAAAGT	CTACAGAGGA	AGATGAGGTC	TGAAATAGAC	AGCTTCTTCC	TTCTTTTACC
15601	TGTGGTATTA	TTCTGTTTTG	TCCTTTTCTC	CATTATCTGT	CTTTCCAGTG	ATGAAATTTT
15661	GATCTGGCCC	TCCCAAGTAT	TAAAAAACAA	GCAAATAAAC	AAATCTCAGT	CATTTTATAT
15721	TAAGATATTG	GCATGCTAAC	TTTTTGCAGG	TTTGTAACAA	GGACCTTTAT	AACTTGACTA
15781	AAAGTTCCTA	AATAAGAATA	TTTACTAGAA	AATTTATTTC	TGCCTGTGGC	CCACATTTGA
15841	GTCAAAATAA	TCAATTAGGA	AAAATGAACT	TGTTTAACTA	AAGTTGACCA	AACTGATCTT
15901	TGACCAAACT	GATCTTTGAG	ACCTATTCAT	CTAAGACAAG	CCAATTAAAT	TCTTGGAGAC
15961	AATTTGTACT	TTAAGGAATT	CTTATAATAT	TTGTAATTAC	CCTCATAACT	TTTTTTTTG
16021	CCCTACTTCT	GTGCTTCTCT	AATATGCAGA	TTATTAAATG	TTGTTACAAA	GCCATTGTCA
16081	AAAAAACAAA	AAACAAAAAA	CTAAACAAAC	TCACATGGTT	AGACTTGCTC	CTTTATGAGA
16141	TATTTTTACC	AAAAATGGAG	GAGTTGAAAA	ACTCTGGTGC	CAGAAATCGT	GAAGACATGG

Figure 8 (Page 5 of 73)

SUBSTITUTE SHEET (RULE 26)

Figure 8 (Page 7 of 73)

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26581	CCAATAAGAG	AGCCAAGAAG	CCGAGAGCGA	CAACTCCTAA	GTCACCAAAG	ACTGCTAAAA
26641	AGGCTAAAGG	AGCCAAGGGT	AAGCAACAGC	AGAAGAGCCC	AACTGTTAGG	AGCGGGAGAA
26701	AGTCAAAATT	GACCCAACAT	CATGAAGTTA	ATGTTAGAAA	AGTGAAGGCA	AGGGCTTCGA
26761	GAGCTTTCCG	GGAGGCCAAT	TTGGAAAGAA	CCCAAAGGCT	GGCCACATCT	AAGAAGTAAA
26821	TATTTTAAGA	TGGCGTAACA	CTGGAAACAA	GTTTCTGTGA	CTTTTAAGAG	CCACCCACAT
26881	TTGTGATGCA	GCTGAGTTGA	AAAGGCTTGA	GATTGGAGAA	CAGTTATCTA	TAGGTTTAAG
26941	AGACCATCCT	GGGCAACATA	GCCAGACTAC	CATCTATACC	TTAATTCAGG	CCAGGCTTCA
27001	CCACCGACCG	GTAACCGGTC	CCTGTCCATC	GCACGTTATG	AGGGGTCCTC	ATTTCCCCGG
27061	GGGTGAGCGA	ACATTAACCA	ACTGACCTCC	ACCOUNTATE	AATTGAGCCG	CACAGCTGAG
27121	TAGATTCTCA	TAAGCTCAAA	CTCTATTCTC	ACCGCCTGTC	AGGTTAGCTG	CAGCATTAGA
27181	GCTCCTTGTG	ACAATCTAAT	CCCTCATCAT	AATGGCACAT	GCAAGGGATC	TAGGTTTCAG
27241	CATTGCTCCC	AGCCCCTGCA	CCCCCTCCTC	CTGAGGTTGG	AGCAGTTTTA	GTCCGGAAAT
27301	TCTTGTGTCA	AAAAGGTTGG	ACACTA CTCC	CGTGGTATAA	TTGTCTTACA	CAAAACGGTC
27361	GTTGGCACGC	TCCCTTAGTC	CCTCCACCCA	TTTTACAAAA	AAGTAAATTA	GTCAAGCATG
27421	TGCTACCTCA	CTCCAGCCTC	CCTGLACCCA	GGCGTTTAAG	GATACAGTGA	GCTATGATGG
27481	AAAAAAGTTA	AAACAGAAAA	AGGCCTTCTT	AGTCAGACGT	TGTCTCAAAA	CTTAAAAAAA
27541	GGAACTAAAA	AGTCTGATGT	CCAATCCTCA	GTCAGAGACT	GCCGTATATC	TAGAGGTCCA
27601	TACATGTAAG	AGCATCTAAG	TTCTCCAAA	AAAGCTCGAT	GGTGCACTAG	AGGAGGCTTT
27661	ATTTGGCATC	CAAACATAAC	TTCTGGAAAT	GCCAGTGTCA	GGGAAGGGAA	GTGGAGAGCA
27721	CTAGTCTTTC	TGTGGTGTCA	TTCTAACTAT	TTTTTTTTT	TTTAACACAA	GTACTACATT
27781	AGGGATCAAT	AAATAGGAAT	CAACCTCTCC	TGTTTCTTAA	TATGCTATCC	ACTGACTTCA
27841	GTTGTTGTTG	TTGTTGTTTT	TCATCTATTC	CAGAATATGG	ATTAGGGGAG	TTTTTTTGTT
27901	TTCCATTGTG	TGTGACTGAT	ACABATARA	ATTATCCTGT	AGCTGAAATT	TAGAATTTTC
27961	TGGAAATCGT	GCTTGCTTAT	TTCCCNACTA	AATTTGTAGG	TTATAGTTGT	TGCAAGAATC
28021	TTACGGTCAA	GTGGTTTGAT	AATTATTT	CTATTAGGTA	TATCAACAAA	AACACACATA
28081	ATGAATTACT	TTAAGTATCT	TATTTATTA	ATATTATTGG	TCTAATACAA	TTGTAACCCT
28141	AGCATACCGA	AGACTGAAAA	ATTTTAIGAA	AAGAATCTGT	AAGTTTCATC	AGACTACCAG
28201	CCCAATTAGG	TTCTGAATTC	CACCTTTCCTT	TCCAAACCTT	AATGGAAATG	TTGGAGGCTG
28261	AGGTAAACTA	CGTTTCTCTT	TARRESTECTS	AATCACAAAC	TTGTTTTAAC	TCTCAGTCTG
28321	GTATTCTTAC	TGATCATCAT	AAACAGACA	TAGTTTAATT	TTCCTTTGAT	TTTTGATTTA
28381	ATTTCGAGAA	ACTTTGAACA	A A CT C C C CT C	TGCTAATGTT	AGTCTACTTT (GGACCATGGT
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28561	GTATAGATAA	AATCAACCAC	ACACCTATAMI ACACCTTTTCC	GAAAGGTTGT	AAACATTATG	TTTAAATTT
28621	ATGAGTTCGG	AATTACTAGG	ATTOTOON	TTGCTTGGAT (STAATTGCCA T	TTGTTTCCCA
28681	AGCCATTTTG	CCTAAATGCT	GTGCCCAAA	AATATGCCTC /	ACTTGCCTGA (CATAGCAGAG
28741	AGTGAGGATG	AACAACTAGC	CTCTCCCAGCA .	ATGGACTGTC A	ACCAGATTCT (CATCACATAC
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29101	GCTAATTTTT	GTATTTTTTC:	TAGACGCCCCC (TGGGATTACA C	GTGCCCGCC A	ACCACGCCCG
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Figure 8 (Page 9 of 73)

SUBSTITUTE SHEET (RULE 26)

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38941	CTCGATCTCC	TGACCTCGTG	ATCCACCCGC	TTTGGCCTCC	CAAAGTGCTG	GGATTACAGG
39001	CGTGAGCCAC	CGTGCCCGGC	CTACTTCACT	TTCTTCATTT	AAAAAAGAAA	TGGGGATAAT
39061	AGTACCTATC	TCATAGAATT	ATTGTAAGAA	GTGCATGCAG	TAATGCATGT	AAGTAGGTGC
39121	TCAGAAGAGT	CGGACACGAA	GTAAGTGCTT	TTATCATCCT	TATCATAATT	TTCATTATCA
39181	GAACAAGGAG	AGACCAGGTA	GAAAATTATT	GTGATTCTTC	AGGTCTGGAA	TACTAGAGTA
39241	GCATCCCAAA	TGAAGGCACC	ATTAAACTTT	GCAAATCTGT	ATGACACCTT	CATGCCAATT
39301	AGAAAAAACA	CCTCTTCACA	ACCCCTTTCA	AGATATTTGC	CTCCTACCTG	CTAAAAACAC
39361	CCATCATACT	ACCCACAGAT	AGCCATGATG	CTTTTTCTGG	GACAGGTGCC	TCTTCCATTC
39421			AGCTGTGCAA			
39481	AAGGCTTGGT	GACAGATGAG	TTACTGGGTA	ACACAGAGAG	AGGATTCAAA	GGAAAAGTTG
39541	AACGGGTCCA	GAAAATGCAT	AGATACATGT	GTAAAAATCT	GGTAAGGTTA	TGACTAGCCA
39601			TTCTCAGATG			
39661	TAAGGAGTCC	TCTTCCAAAA	ATAGGAAATG	AAATGACATA	GGTGTATGTC	TCTGAGGTGA
39721			TCTAGATGCA			
39781	GAGAGGTCAC	AGCTAGGGAT	CACCGGCATG	CAGGAACTCA	GAAACCTAAA	TGGGGAAATC
39841	TTTTTGAGGA	AATGAACAGA	GAAGGCTAAA	ATCAAGGAGT	TCGTCAGGCA	ATTTCTATGT
39901	TTAGGTTCAA	CTCTCTCCTG	AAACATGAAG	AGCTCATAAA	TGCACTCCCT	CTTTGAGTCT
39961	CTAGTTTTGT	CTCCTTCCCA	CAGTGAGTCT	GCAGGCTGCG	TGTCACTCAC	GTTCAGCTAA
40021						GGCATCACAA
40081			CTTCTCTCTT			
40141	TAGACCTGGG	CACTATTGGA	TTTCAAGAAC	CATTATCTCT	CATCTGGAAA	TGCTTATTGG
40201						ACCATATAAG
40261						CATAACCCAA
40321						CTTTTCCCTA
40381						CGCAAACTCA
40441			TTTCTAATCT			
40501			GACCTCCCAT			
40561			GGCTATCTTG			
40621			TTCCTGCCTG			
40681			TCATTCCCTA			
40741			CCTGTACTAC			
40801			TAGAGACTTT			
40861			CCCAAAGAAT			
40921			TCCCACTGTG			
40981			ACCCTGCCTC			
41041			AGCAATTCCA			
41101			AAAAGCAAGT			
41161			CTCATCTCAC			
41221			CAGCTCCTGG			
41281			GGGTGCTCCA			
41341			CAGCCCTGGT			
41401			TCCATTGGCT			
41461			GCCCGACACC			
41521			GGCACTTGAC			
41581			ACTGGAACTT			
41641			CAAGCTACAG			
41701			GGAGAGCAGT			
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42061			AGGCTGGAGT			
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Figure 8 (Page 13 of 73)

Figure 8 (Page 15 of 73)

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51901	TAGACTACTC	TGGCCAACAT	CCTCAAACCC	CATOMOMA	TCACTTGAGG	TCAGGAGTTC
51961	TAGCCGGGCA	TOGTOCOTOT	ACTOCOLOGO	CATCTCTACT	AAAAACAAAA	AATGTTATCC
52021	GAACCCCCCA	TGGTGCCTGT	AGICCCAGCI	ACTCAGGAGG	CTGAGGCAGG	AGAATTGCTT
52081	GAGAGAGGA	GGTGGAGGTT	TRARRACT	GAGATCACGC	CACTGCACTC	TAGCCTTGGT
52141	DAGAGAGAAA	GACTTGGTCT	CCCCAAAGAG	AAAAGAAAAA	TGAAATTTCA	GCATTATAGA
52201	CTCTTTCCC	TTCCCCTTCC	CCCCAAACTT	TAAAAAAGCA	GAAGTCTGCA	TCATAAAATG
52261	CARTTCTCAC	ATGTTATTTT	TATTATAACA	AAGGAATCTT	GCAAGGCTAC	CAGATCTCAG
52321	CAMITGICAC	TATGTTCTGT	AAAAATCACT	TCCTAAAATG	TCTGAATTGA	CTGCTTGTCT
52381	CTCCATTTTC	TTTCTCGTGT	CATACTGCAA	TGGATATCTG	TCTTGTTAGT	ATAAATATTT
		TTGTTGTTAA				
52441		ATGTTTAACA				
52501		TTAAAGAAGG				
52561	CTCCAAAAAC	CGAGCTCCCT	GAGTGAGCAA	TTCCTGTCCC	TTTTAAGGGC	TTGCAACTCT
52621	AAGGGGGTCT	GTGTGAGAGG	GTCATGATCG	ACTGAGCAAG	TGGGGGTATG	TGACTGGCAG
52681	CTGCATGCAC	CAGTAATCAG	AACAGAACAG	GGATTTTCAC	AGTGTTTTTC	CACACAATGT
52741		TAGATAACAT				
52801		AGGCTGTCTG				
52861		GGCAGAAATT				
52921		AGAATCTCAC				
52981	TCTTCTTGAA	AGACAGATTG	ATAATGATTC	ATATAGTACA	CTTGTGCTGA	AGCATTTTGG
53041	TGAGCTAAGG	TAGTGATGAA	GCTTTTTATC	ATTTGGAGAA	GTACAGGTAG	CAAACAAGGA
53101	AĞCAGTAAGC	AGGTTTCTAT	TAATATTATA	ACTCCTATTA	TAAGAGTTTT	AAATCTTCTT
53161	AGCACTCGGA	ACCATTTTTC	AAACATGGCC	CCAGAAACAA	ATCCATACCA	CACCTACATG
53221	GGCACATGTG	CCACTTTTGT	CATATTTCTA	ACTATGTCTT	CAACTACTTG	CCCTTAATCA
53281	TCTATGTGTA	GACAGCAATT	AGTAAGGTTA	AATTTCCTAC	AGACCCCTCC	TTCAGTTGCT
53341	AGCAAGTAGT	CGAGAGCCAA	TCCATTTTGA	TAGATAGCAT	TTTGCATCTG	AGTTTCTTGC
53401	CAGGCCACAG	TAGTCAGGGC	TCTGCTGGTC	TTATTAGTAA	TTATTTCTAA	GACAGCTTGT
53461	AACCGTATGA	TTCAGTTGAG	CATGTAAATG	GGGGTCCCAT	ATCCCCACAA	GCCGTCTTGT
53521		CAGGCCCATA				
53581		TAGCTATGCT				
53641		CAGGGAAGCC				
53701		TGTCAATAAC				
53761		ATATCCAGTA				
53821		CAGTTTGCAA				
53881		AGGTGGCTGT				
53941		CAGGAAGGGT				
54001		TTAGGACCCA				
54061		CTGTAGGTAC				
54121		ATACATACAT				
54181		AACAAATTTC				
54241		GGTTTGAAAT				
54301		TCAAGGATCC				
54361		AGAATCAAGG				
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Figure 8 (Page 17 of 73)

Figure 8 (Page 19 of 73)

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TODAAAADAD	TTTTTDDATA	ADATTADTOO	ATOOTTTOTA	SSSTTSSSS	ACCCGCCAC	17809
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CACACACCACC	CACACACACA	TACACACACA	ADAATDOOTA	ACGTGGAAGA	CCACATAGTT	τ0909
				DAADTAAAAT		17509
ADADTDADTT	DOTATABTOA	DTODDDTDTD	AAADDADADT	CCCATAAAGG	TATADAAADT	18709
DTADTATAAA	DOTTTTATAA	DDDTDDJTDA	DDDDTDADDD	TASTSSTSAS	ATACTCCTATA	12709
DDADTDADAD	DTATADDTDT	DOTTATDACT	DETAATAADT	DDADADATTA	AADAATTADD	T9E09
ADADAAAATT	TADDITITOD	TOTOTAOTOA	AATDDAAADA	AADTAATDDT	TOADBAAADD	T0E09
TAAAADDADA	TADATTTAAA	DAADAADTAA	ATAAADTATA	TATATATATA	TATATATATA	17209
TATATATATA	TATATATAT	AAAAAAAAA	AAAASTSTET	TOTOAGAGTOT	DADADTDDDT	18109
SSEASSTOAS	STTADDADAD	TADADTDDAD	TDADDTTDDA	CAGGAGGCAG	TDAAATTDAD	12109
				STESTSSTS		19009
ATTAAAAAAA	ATAAAAAADA	TOTOTAOOOO	AAA DTDTTAD	AADDEDTDEE	ADDADADTTT	10009
DADDADTADD	SPECCEGEC	DDDADDDTTT	DECCAGOGO	CATGCCAGTA	TODDITODOO	T 7 6 6 S
DDDDDDADT	ATDTAAAAA	TADADDTDAA	ADADAAAADA	GCCAAGAAGA	DAADADDTAA	T8865
DATTAATDAD	TADDAAADDA	DADTAADTOT	DTAAATTAA Đ	ADATODTADO	AATDTDDTDA	TZ865
DADAATTDAA	ATABBTADDD	TADDAADTAA	DTTDDDDDTTA	ADAATTAAAA	TGTTTGTCTC	19465
TTAAATAƏTT	AAADAAAAA	DADADADAAA	CAGAAAAAA	AAACCACACA	AADAAAAADT	10465
DIAMADADID	ATACTEGATO	TAAATAĐĐAA	ADTATTĐAAĐ	AABABABAA	DDDTAAAADD	17965
ADDTDADDTD	DADTDDATAT	ADADADADTO	TODDAATODA	TATADADDTA	TATATOTOTT	T8565
TDAAATDDAT	ATOTATADAD	DTDTDDADTD	TATTTATADA	TATATADDDT	CCAATATCTC	TZS65
DATATATAA	ADDDDTDDDD	TOOADTOOAT	ATADADDTDT	ATADDDDDTDD	COTOTOTOT	19765
DDADADDADD	TOADTDDDAD	AADDADAADD	DAADADDDDT	CACCGTTACC	DODIDADIDI	T0#65
SSSASAASSA	DDDAADAATA	ATAATODDDD	ADDDATDAAA	TAGAGACCT	DADDAATAAA	T 7 E 6 S
AADTOTOTTA	AAATTƏAƏTA	TTCATTAGTT	AAAACATTTG	DDDDTDTTT	TATADAAATD	18265
Deparater	TADTAAAAT	AAAADATA DD	DTATTDAAAT	CAAATTATAA	DOTTAAATDA	29227
ADDDDATDDT	DAAADTTTTA	AAADAAATTT	AATTOOOTOO	TOTODTODO	GACAATATGA	T9T6S
AATADDDDTT	AAADADDDAD	TCTTTTTG	TTTTATTT	TAATTTTDDD	TOTTTADTTT	TOTES
ようらすらずつつらす	TT9T799992	DTDDDDTDDD	ACCAGACCCA	ATTTSTABST	AGG1/CAAAGG	T 7065
TTOTOTAATA	DAATADATAT	STAABBTSTB	TCCATACAAT	TTTDDTAADA	DTTTTABBBA	18685
2428A2AABA	AGABCAGAAC	DTAATDADTA	DDTADDTDDD	ACGTGACTGG	AGCAGCGGGT	7768S
ADDADTTADT	TADTDDTDDD	CTGCATGAGA	TDDDDDAATTO	TOAACATTOO	SSAATT LTDD	T9885
PTSTDOTTAA	DDADTDADTD	DOTOBADOAA	AAADDTDTDT	DTDDADTAAT	LECTROPHY	T0885
エエつつむてつむむむ	DAATCODOTO	AATDDDDADD	DDADADDTDD	TABBBATBDD	CT PPPPTTCC	17785
SOTTARACTO	DOTOTOTA	DACOTOTODO	DADDDTDDAT	SGCAATTGCC	9500V001.t.e	T8985
Desagraga	TOSTOSTOOA	AGGGCCTGCC	DOADOTOOTA	CACTGCGTTG	SSASTIBLES	17985
つつつすすつつすする	ADDDTDAAAD	AADABTDDD	TATAADDTTT	TODIAATTOO	DDAADDATAU	T9585
THE THINK ADDAA	TTDDDTJATT	CATCCGCGGG	DDDDTTTADD	CAGATTGAAT	₹₽ 35335££	10585
ADDAAADDAT	TOTOTITI	DATTTTTTA	TTDAAAATDT	CAGGTTTCC	AADADTD016	77785
T0040T0022	AADTOTTOAT	ADDATTOOTT	TTTTTTT	TOTTOTAOTA	AATTƏTAATT	78285
SOTEASTITT	AADADAATTO	DDDDDTDDAA	CCTGTTAGGA	DOTABATTTA	DAATTAATDD	28327

64801	GCCAAGCAGC	AATGGCAGGT	AGTACACACA	CAAGAGGCAG	ATGATACAAC	A C A TCCTTCC
64861			CCCCACAATC			
64921			TCCATACAGA			
64981			TATTATGACA			
65041			TTACTATTGG			
65101	GAACCCATAG	DACADATAAT	CTACCAAATT	TTTAACATTC	ATTTTCCCA	ACCOMMENCE
65161			GAAAACTTAT			
65221			ACTAATGAAC			
65281	AATCTTAAGA	CARTACTTA	TATGGCAAAT	CTTA & CTTCT	TAAATITIT	AATTCTTAAA
65341			AGTTGAGTTA			
65401						
65461			TTACAATCAC			
65521			TGTTACAAAA			
65581			CCTGAGTTCT			
65641			AGTTTTCCAA			
			AAACCTGCAA			
65701			GTCTATGAAT			
65761			TTTAGTCACC			
65821			ACAAACTCAG			
65881			TTTCACTGAA			
65941			AAACGTGTCA			
66001			TCCAAAAGTT			
66061			GAACCTCTAG			
56121			CTAAGCTGGG			
66181	GCAGTGAGCT	ATGATTATGC	CACTGCACTC	CAGCCTGGGC	AACAATGCAA	AATCCTGTCT
66241	CAAAAACAAA	AACAAAAAAC	AAATTGCCTA	TGCTGTGGTT	ATCTCACAAT	TAATAAAAAG
66301	GAAAAAAAA	GTATGCAGTC	TTTGTAGGTC	CTTGGGGTTT	GTTGGAACTC	AGAAAACAAT
6 636 1	ACCCCAAAAT	AAAGACCGCA	GAAGCCAAAG	TTTTTCTCTG	ATCTTCTCCT	GCCCTCCTGT
66421	CTCTGAGTCC	CATTCTCCCC	GGAGTCTAGC	CATAGAAATG	AGAATTCCTC	TTCCTCAAGT
66481	TAGGTCATAG	AAATCAAAAC	ACCTTTTCCC	CAGAGCCCAG	CCATAAAACC	TTATAAAAAT
66541	ACTCTAACTT	TCCCTCTGTT	TTTCTGTGTA	AAAACTGGCC	ATAAAGAAAT	TATCTGAACT
66601	ACCTTATTTG	ATCATAGATC	ACCAGACCGC	ATTCCAGAGA	GGATCCAGAA	GGAAGGAATG
66661	CTGCACAGAG	AGGCGAAGAA	GAATCTAGAC	AGACAGGCCT	TGCTGGGTTT	CCCTACTCTG
66721	TTTATTAGCA	ATCCTATTTC	TACACGGCGG	CCCATACTTT	GTTGAATCTA	AAAATAAAA
66781	ATGGACAATT	TCCCCTGTAC	ATGTTAATAC	ACATTAATAA	ATTGGATATA	AATTGGATAA
66841	TTTATTAATA	TACACATTAA	TAAATTGGAT	GCAGCCGGGT	GCAATGGCTC	ACGCCTGTAA
66901	TCCCAGCACT	TTGGGAGCTG	AGGCGGGCAG	ACCACGAGGT	CAAGACCACC	CTAGCCGAAA
66961	TGGTGAAACC	CCGTCTCTAT	TAAAAATACA	AAAGTTAGCT	GGGCGTGGTG	GCACATGCCT
67021	GTAGTCCCAG	CTACTGGGGA	GGCTGAGGCA	GGAGAATTGC	TTGAACTCGG	GAGGCGGAGG
67081	TTGCAGTGAG	CCGAGATTGC	GCCACTGCAC	TCCAGCCTGG	TGACAGAGTG	AGACTCCGTC
67141			ATAATAATAA			
67201	TCCTATTAAT	CTTCCTCTTG	TCGGTGGTTT	TCAGCGACTC	TTCAGAGGCC	AAAGAGTAAG
67261			GTTCTTATGT			
67321			GATTATTTCT			
67381			ACACTGACCC			
67441			TCTTGCTCTG			
67501			CCTCCCGGAT			
67561			CCACCACCAA			
67621			CCAGGATGGT			
67681			GGATTACAGG			
67741			TCTCCCTTTT			
67801			AATTGTTTTG			
67861			ATTCTCTTTA			
67921			GCTCCAGGGT			
67981			ATAATTATTT			
	CICACIGICA	WILDSOLL	UTUVITUTI T	TATCCACCA	THOMCICIO	1710104410

Figure 8 (Page 21 of 73)

Etgure 8 (Page 23 of 73)

DTDTTDTDA:	CACGTAGGA	r ttjattðaða	STDAATTADA	AAŢĐTAAAAŢ	T AADDDADDTD	T988L
TDATADATT.	r toattteta	4 DTDAADATO1	TDAAATTATI	TAATTATTO	TAAAADADTA	ፒዐ ፇ፝፝ቝረ
TOTOTADOT	T TADTTAADA	TTATATOTTI	TTTADTDDDA	TITIOAATA	TAADATAAD	14341
DATTATTDA	D TAATTTƏTT	D AAATATADAT	TAAAADDDAA	DDADAGOTT	ATATOTAGOA	18297
AADDDADTD	D TTTATDTATA	AAATAAĐAĐE	DOATADAADD	AAADTƏƏATI	AADADTAADA	14221
DDAADDADD/	A ADTAATTAD?	r taatttadta	ADTAADTDTD	TATETETTO	ASTSATATAT	T9T%4
ATDATDTAT/	4 TADADTDAD1	r adaaaaatte	CATTCAGGAA	AADATDAAA	WALCAMGIGI.	TOTEL
DAADTTAAA	O DESATIBLES	ADADTTADTO	TTDDAAATDT	TODDTAADAG	T.T.J.J.J.J. (10006
TOTATOTIT:	D DAAAADTDA1	r ADTTDADDTA	TODITEDATI	ATABADDAAD	CATEGRACAC	18657
DTTTDADAT	4 AADADTTTA/	AADTTTATTC	CTTGTTCAAT	DDACACAGG	TITLEGILES	13921
TTTTDA9AT/	4 ATOTTADE	CCCGGCCAGA C	AGCCACCCCG	DTDDDDADAT	TABBBTTBT	13867
AAATOOTOOS) ADTODADODE) TOTABLABIES	DADTOOTOAA	ATTOTOOTO	TellectCAG (73867
ADDADTTTAA	A DODADADAAS	ATTTTTATOT	YLLLTYTT Y	DODDIADDAT	MANTECACGC (T\$4.57
DATTABBBT?	DATDAADDTT	CODEADTOODI	DOTOTTABTE	AADTTBBBTU	Cettcecctc	18987
AADSTDADTD	D DADTOTAADE	DESCAGIGGC	CCAGGCTGGA	DADTOTOR	AT MEMPTICAL C	12957
AASTTTTTTT	TTTOTTADTE	DABBITTOODI	ATCTGAACTT	ATTOOTAGAG	CIPPPROPER	T9582
SOTTTTO AD'	r ADABBABTTS) TTTD55T5TT	CTCAATTTGT	DYDTDTDAA T	TULLULL	T05EL
TADDTTATA1	AADADATAAE) AAAATAAATD	TOTOTOTAOT	ATTTATAATA	TAPTCAACCT	19457
SAAAASAAA E) TADSSTSTSA	BAADTDDDAD	ACCETTICCA	SSATSTITIO	AATTTTADD	IBEEL
TOTODITITIA	TGGACTTGAC !	TACTGCTGTT	TACCCCAT	TOTOTOTORS	TTGGCTGTT	
DTATATAAD1	TTOOTATOAC	TOTOOTOAAO	AATOOTOTOO	ADATTOOTIA	DTDDDADTDT	73321
DDADDAAAAA	DAAADTDAAA	AASSAASATS	ATADDUTTAA	TOTOPPARA	DAAAAADTAT	13267
STSTSTSSS	ADADTDATAD	AASSASTSSS	ADDABABTIT	DADAAJJa14	STATCACTTG	73201
ADDADDDADT	DDDADDDTTT	DAAAADTDTA	ATATODADAD	Linging	DDDTDDDAAT DTTD4DT4DD	19167
DTDATDDDAA	DADADTADAA	AADDTADATD	DATTUDATTA	TAALOALLIA	222772244T	13087
DDTDTTDAAA	DITADIAAAT	TTOĐADAAAA	AABBADTBAB	TOOLIAMILA	DITIONATATION	73027
TATTƏTƏƏƏT	DODDIDADOD	TOADDDADDD	AUTTIOTOAT	TORORAL DAD	AAプロス1カロホン つますつのも生まるT	15627
DAADTDTDDT	TOTTATOTO	DTDADDTDDT	ATAAAADTOO	TO TO TAIL TO	CAGATAGGAA	72901
ACAGACAAGA	ADDDADDAA	DTDDDDDTAT	TITOAAAAAAA	SIMMERTANT	6676777779	12841
DIDAAAADDA	DTATAAADTT	ADAAADTTDD	STITALLE	TOTING TOTAL	อออนานาออน	18727
DDATTTATAA	ASSASTSSAS	TTOOODADTO	THACTITUMAT	TOTT A ATOTA	ATTTE	12727
TOOODOADIT	TTADODAADD	TOTTTTOTTO	TOTTIAIAI	ACAGATGCAC	TAADDDATTD	19977
TATTATTAT	DADTAATDDA	SSSTGCACCC	COTACCOLOR	DATTANDED	られるインシエンシ	10927
DTADTDDDAD	TOODIOOTOO	TADTDAAAAD	AACTGCCCCAMGG	CGGGAACTACA	DATDADATDA	18527
TADTOTTOTO	AASTSASATT	DDADDDTTD	A CT TOCTOCT	ADATA 475T	PTPTPBATAD	18727
DTATTTATAT	TTTAAATTTT	TATATTTATT	TATAIAIAI	TOTIGGRERER	Trymp ATTTAT	12421
TATAADDTTT	TATATAAATO	DATTDTAAAA	DITIMADED	エンイエイイエンエエ	TTOTTTOTTA	19827
ATOOTITEAD	DOADTAATTO	TODATADAAT	AMUMMOT 1 DO	ACTATTAGOA	TATTOADTOT	10827
DOTAMADIDI	TTOADTTTDD	ATDDDADTIT	COMMITTIES	AASAAASTTT	ADTADADTIT	12241
DTDTAAAAA T	DTDDAADADT	TABADADADA	DDEATTOTTA	4424277442	ADTODIATAD	18127
ATATACGTTT	ATOOUTTATO	ADAADTADTO	TIMITORIST	TOAATATTTO	STSTTSTASS	12127
TAADOTTOTT	TOTACOTITIA	DOTTTAAADT	TELLAMICIAL	ATTRACTOR	STTADTSATA	19027
TAMAMAAT	CACAGCACAG	AAAATƏTƏT	ACTANTAMO	ADTESTAGTAG	AAAATTADAD	10027
DIDDIDITITA	TTATTTTDA	TTOTABTODD	ATJULTANA	TOAADAATTD	STSTSTSTS	t \$6TL
TGTCCCACCA	STATSTST	TTATATTOOT	AUTATOTAC	ASSASASTTO	TTĐẠATATĐĐ	TBBTL
DIDIDDDTDA	TTGTGCTCTC	TTTTTTTT	CATTIAIGE	ハコーナー・インコー・インコー・インコー・インコー・インコー・インコー・インコー・インコ	TATOTTTTOT	17877
TTGAAGGCAA	DATAAAT DD'I	OADATODAOO		Agrrana	DODATTTDDA	T9LTL
ATDADDAADA	ATTTOTAAAT	STATABTOST	OFFITT ATTOT	コヤンエインエライブ	DIDDDATITI	TOLTL
ADDITITITA	TAAAADDATT	TATATOODTA	PATROTORM	BABBATTTTT	DATADTOODD	T # 9 T L
TTAATAAATT	AATAAATATA	ATTATAAATA	AATADTDAAA	OUTTITITIE	DDADTDATAA	TBSTL
ATAAATAAAT	ATTADUTATA	TADADTAATA	A A DIT ATTENDA	A ATTTTATAA	TATTTAATT	TZSTL
TATATOOTAT	STAAATASAA	TATATAATOT	ポイポング 453 T. A. T. T. A. A. S. T. A. T. T. A. A. S. T. A. T. T. A. T.	ANATOBOTAT	ATAAATAƏTA	19 5 1
DITITIONADO	DOTAADOUAL	TOUDIAMOII	*************************************	CCATTTTTTC	TTTDATTTDD	TOPTL
AADATTTTDA	Seekasirkak	TODOTA	プロペントンンパンニ	nabbbTDD8A	TTADADTTTD	17877
		TOTOTOTOT	つうなつてつむなうつ	ATDADADDIT	STEETCTGCTC	11281

77761	CGGCGCTTTC	CCACTTCTAC		CATTCTCAAC	ATGTCCGAGA	CTC
77821	CCCTCCCCC	CCCCCCCCCCC	CTCCCCACAA	CCCCCCCCCC	AAGAAGAAGG	CTGCTCCTGC
77881	CCCTCCCCCT	ACCOCCCCTC	* CCCCGGAGAA	GGCCCCIGIA	AAGAAGAAGG	CGGCCAAAAA
77941	TCTCCCCCCC	ACGCCICGIA	AGGCGTCTGG	TCCCCCGGTG	TCAGAGCTCA	TCACCAAGGC
78001	TCCCCCCCC	TOTALAGAGE	GTAGCGGAGT	TTCTCTGGCT	GCTCTGAAAA	AAGCGTTGGC
78061	CCTCACCAAC	CCCACTOTICS	AGAAAAACAA	CAGCCGTATC	AAACTTGGTC	TCAAGAGCCT
78121	CARCARCARG	GGCACTCTGG	TGCAAACGAA	AGGCACCGGT	GCTTCTGGCT	CCTTTAAACT
78181	CAACAAGAAG	GCAGCCTCCG	GGGAAGCCAA	GCCCAAGGTT	AAAAAGGCGG	GCGGAACCAA
	CARCARGARG	CCAGTTGGGG	CAGCCAAGAA	GCCCAAGAAG	GCGGCTGGCG	GCGCAACTCC
78241	GAAGAAGAGC	GCTAAGAAAA	CACCGAAGAA	AGCGAAGAAG	CCGGCCGCGG	CCACTGTAAC
78301					AAGCCCAAGA	
78361	AAGTGCTGCT	AAGGCTGTGA	AGCCCAAGGC	CGCTAAGCCC	AAGGTTGTCA	AGCCTAAGAA
78421	GGCGGCGCCC	AAGAAGAAAT	AGGCGAACGC	CTACTTCTAA	AACCCAAAAG	GCTCTTTTCA
78481					TTACTATCTG	
78541					GAGGTATCAG	
78601					TAAACCTTTA	
78661					TTGGGATGGC	
78721					AGTTTCTTAG	
78781					GCTCCCGGGT	
78841					GGAAGACGTT	
78901					TCTCTTTATA	
78961					TGTCTCCTGG	
79021	AACATCCAGC	CCTGGGAGGA	GAGTGCGTGC	AGGTACCTTT	GTCCTACATT	CCTCTGCTGT
79081	TAATTTCTCA	TTCCTGTGGC	AACGAAGGAA	TGCATTTAAA	AAACAGCCAC	AACAGCGGCA
79141	ATAGCCCTTC	CTCCACCCAA	GGCAATCGTG	GACCTAGGGA	GTTTTTTGTG	CCACATAACA
79201	TGTAGCCTTC	CGCTAAACTG	ACAGGTTTGA	GCGTATCGAT	TTTGAGCGTA	TCGAAAGCAC
79261	AACTTTTAGC	CAGCCATTTT	GTCCTCGCAT	GACTACGGTT	GCTTATCCTG	TTTAGACAGA
79321	CAGCAACATT	TAAAAATCGA	AGTTCCTTTA	AACGTATTTT	GTTTGGCAGT	CCAAATGTTT
79381	CTATGCAGAA	AACAGTATTT	GTACTATTAA	CTATGAAGAG	TGTATGGATA	AATGGGAGAC
79441	ATTTCTAATA	AAGGCCTTCG	TTAATGGTTC	CCTCTGTTTG	ACATCCATGG	TGCTTCTGAA
79501					CTGGTGGGCA	
79561					GCTCAATTAT	
79621	CCACAATGAT	TAATATAGTG	AGTTGATTTG	TTAGTGATAG	TGACCACGGA	TTCATCCCAA
79681	GAAAGAGAAA	GGGGAGGGAG	GCAAGCAGAG	AGACAGGAAG	ACAGAGGCAG	GGAAGAAGGA
79741					ATTTTACATT	
79801					TATGGACATA	
79861					TCCCGTGCCA	
79921					ATAATTTCAT	
79981					TCATTCCCAC	
80041					CTCAATGCTG	
80101					GTATCTTATC	
80161		•			ATTACTGGCA	
80221					GATCACAGTC	
80281					TCTGTCACCC	
80341					GGGTTCAAGC	
80401					CCATGCCCGG	
80461					TGGTCTCGAA	
80521					TACAGGCGTG	
80581					TTGTTGTTTT	_
80641					CATCATTGCC	
80701					TTTCAGGCAT	
80761					GGGAGACTAA	
80821					TCTCATTACT	
80881					TATATTAAGA	
80941					AAAAAAAAAG	
30341	C1CC1CC1G1	TICICCAICI	CANCATCAAA	CAATTAAAAA	DHAMMANA	301000000

Figure 8 (Page 25 of 73)

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SSACATO AA	A ADTOADTTA	r DADADTAADE) TABAADTOTA	DDTTTAATA/	AADATTADAA	87421
つなわなてなてつつん	1 TTADOTTTA	r ottaabttas	AAADDADDID	TCAGCCACC	ACATATTATC 3	19578
AAADAADTA1	r TOAATOTTO	4 TAAADTDAAA	, DTTTTDTDTA	ATAADDATI'A	CTTMCTTGAC 1	10578
AAAATTTA	ADAAAATT	r AAADDTTTTA	(DDADADAATT	CAAAGGAGC .	DTADATTDAA	87241
DATTTDAAA	ATTOTTAADA	A AASSTOTSAT	SOBAADADDDA	DATATODAAA	DATDADADDT	TBTLB
DATITUTOTOAS	D TEABASTDAT	D BADDDTTADA	L DAAAADTTAT	SACATOTADAS	WWW.I.C.I.LV.I.	TZTLB
TABATOTEE	1 DETABAADE	D TOOTTAAATO	DADTDAATAA	DDTDADTDD1	DATCOTTOTA	19048
ABBAADTTDE) DTAADAAAA	A AAAAAAAAAA	CICCGICICA	ADABDBADAL	YEVESS TONE	10078
4つつTOADDT2	ADDBDDTTA	ABDDBABTBA	CGGAGCTTGC	25A5557777A	WELFCCATER !	T 7698
o4554755A5) TODDADDOTT	r DATDBADDDT	• DATOTOODO	STRILLEGGGG	T TWPCCPBBC C	T8898
AAAAAADATA	(AAAATDATD1	r ttddddddaaa	TAACACGGTG	Septitotrace	AUAUU MUMU	12898
947T99A97 A	LOTABBOBBB	S ADADOODDAD	AGCACTTTGG	TOOTAATOTO	Teer to Voce	T9498
ออวอวออออ	D DDADDAADTT	r Deetaabaat	DOADTOTAKO	CerriteceAcc	ATMOTTOWIN.	T0758
TOTADTADTE) TATADƏTTTA	§ DTDDATTDAA	TAATAĐĐĐĐT	ATDATOTATO	WATTCHWCHT.	T\$998
TTTDAAADAE) AATDADTTT#	ADSTTTATAA	AASTASTOOT	AADATTDTAT	Attercievic	T8598
2242T4444T	· TATTTTOOI	CACTCAATCC	CTGCTATTTT	PERCETTCAGG	SOCKLESS T	TZ598
エコタエコシタダエゴ	* ATDADDTTTA	LAAATTTTA DT	TCTCTCTCTC	DIDIDIDI	PATIBIBITAG	
ADATDDDDD	DABABADDTS	DAADOTTOAT	DTOODDAAT	STSAACACAA	AUIMAIJUAA SIMMOMOMAA	19898
DTDDTATTTA	TCTGGGGCAG	ASTSASSET	TTAAAATƏJA	2222442444	ANGCENTAGE	T0798
DIDADATAAD	TTTATOTADA	SADSTOOTOS	Teeschere	AAULULIAAL	#22#655##2	17598
TADTOOTOAD	AAADDTADTA	DEAAADOTAD	SASSASSASA	ADADIDITA	CCCAGAGGAG	18298
DOTOTITODA	ATDABTDDTD	TTATATTTOA	DITIODAATITO	OIICAMAIIG	TEACHTON TOTAL	86221
AATTOAOTOA	ADDDADDDTE	TTDADADDTT	SSASAAATT	DAMA I MANJA	TOARDTTATO	19198
TODAAOAAAA	TOTOTOODT	TODADATOTT	COLUMNIATION	DESTRUCTE	T4427TT4T44	T0T98
TATDATATTT	DATATTTDAT	TDADADTAAD	TATOASARAS	T 4000000110	STAASTAAT	17098
TOADDAADDA	DADTTAATAA	DADATTTTAA	DAMI JAMAT J	#2222222TT	4 エンシエンエダンエ	18658
DTDADADADD	TAAAATƏTƏT	TOADOTODAA	CTTGAGGTTG	DTA ADATTOD	STITTAGTADA	82927
TAGAATDAT	DDTODIOTAT	ADTAADDETD	2522256252	TTAAASTSAT	TTAADDDDTT	19858
TATTOTATTO	DDATTTTAAD	TAATADTATA	CTTO AD ADTO	TOAGATTATA	DAADATTADT	T0858
ATADTATATA	TOAAASSAAA	DAAADTOTTT	TATT TO T	ADADTATATA	TOAAAOTATO	14728
GACACCACCA	CTTAGGCCAG	ATAADAADDT	TACCACCACCA	45T222422T	ATTOOTOTOT	T8958
TTDADDTDTD	ADTITIOADDO	AAADDTTTTT	TASSASATTA	atata and a	TADADTDDAD	82621
TOTADAATDD	ATTSSSTSAS	DTADATTDDA	TOTATTOADT	DTTTTAATAA	TOTOAOTITO	19558
TOAAADDDTA	ATTDATTDAA	ADDDTDDTDD	PATTER SEPT	TOBACABBAA	AADTAADDAD	10228
SCTGTTATTG	DDAAADT'''T	TOTTTOOODA	AATDAAATAT	SAAAADSAAA	ATTTOTTTT	T\$\$58
DOTDATTOTT	TAATOSSTOI	TOTAATATTT	Cheechan	POSTERIOR	ADDAAAADDT	18258
ATOTTOTAAD	A'I'I'I'DOADII	TOTALTT	AATADDAAAT	DOTOADTADD	TODADOTTOD	17858
ADTTADDAAD	TODATAMAN	TAASSASSAS	POTTATTTA	DTDATAAAT	TITAGGTIT	19258
TOTATTTDAD	DIAMINAMO	TCAGGGTCTG	TOTOTTADDI	AATODAOODT	DDTDDDTATA	8250 <i>T</i>
Detettetec	TOTOTARETOD	ADAADTDADD	5557555575	PTDADDDTTT	TTDADTAAAT	T \$T58
DOCTOADTDA	COMPONENTS	DADDDATATT	AAADTAĐAĐA	OTDDDTTDDT	TODITOTOAD	T8058
TOSTOTOROLL	AAACICAAA	TTADTTTAAT	ASSAATITIT	DAAADDADDD	DTTTAADADT	12028
TTAADDDDAD	ADAADIAIII	TTATTTDAD	ATDABABBAD	ATODODADOD	ADDAATAADD	T96#8
ADDAATUTIA	COLLIGICA	ACAGAGCTGA	9AT22TT2A2	DTDTTDDTDT	SECENTICO	T0678
OTTOTODADI	TITITION OF	TTATTTATA	TTTDATAATD	TOTOTOATOO	ADDDAAADTD	T 78 78
AADDAADADA	DETERMINED AND A	AAADAADOTT	TTOTADADOT	AADADDADDT	DTDDTTDATA	18728
TADADTDATA	Attonomical Cartesian Attonomical Attonomi	ATATTTTDDA	DADDADADDT	TOTATOAAOO	DTDATTDDTA	1272B
- 10 1000 4774	イエインコース・インプ	TTTADADDDA	TOADULTION	THT I WIND OU	WUTTALL	T99+8
tacacaca	TOTALABOTIT	DTTDATDATT	J.I.V.J.I.O.I.I.I.O.	WITTIIONOT	TUTURALLA	T09\$8
		TTAAAAADDA	AT-CEAST-FEE	WALLING INC.	* * * * * * * * * * * * * * * * * * * *	T7578
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PTAPAAATDT	DIDITITOODA	DADTTTAADT	DITODITODOI	DTATDTDDAT	TADAATTTTA	T 7 Z 7 8

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90721	CCCCGACACC	GGCATCTCAT	CCAAGGCCAT	GGGGATCATG	AATTCCTTCG	TCAACGACAT
90781	CTTCGAGCGC	ATCGCGGGCG	AGGCTTCTCG	CCTGGCTCAC	TACAATAAGC	GCTCGACCAT
90841	CACCTCCAGG	GAGATTCAGA	CGGCTGTGCG	CCTGCTGCTG	CCTGGGGAGC	TGGCTAAGCA
90901	TGCTGTGTCC	GAGGGCACTA	AGGCAGTTAC	CAAGTACACT	AGCTCTAAAT	AAGTGCTTAT
90961	GTAAGCACTT	CCAAACCCAA	AGGCTCTTTT	CAGAGCCACC	TACTTTGTCA	CAAGGAGAGC
91021	TATAACCACA	. ATTTCTTAAG	GTGGTGCTGC	TGCTATTCTG	TTTCAGTTCT	AGAGGATCAA
91081	CTGGAATGTT	' AGCGAAGACA	AGTTTTAGAG	CCAAGGTTAA	CTTGGACGGG	GCCGTGCGCG
91141	GTGCCTCTTG	CCTTTAATCC	CGGCAATTTG	GGAGGCCGAG	GCGGGCGGAT	CACGAGGTCA
91201	GGAGATGGAG	ACCATCCTGC	TTAACACGAT	GAAACCCCGT	CTCTACTAAA	TARAGOATAA
91261	AATTAGCTGG	GCGTGATGGT	GGGCGCCTGT	AGTCCCAGCT	ACTCGGGAGG	CTGAGGCAGG
91321	AGAATGGCGT	GAACGCGGGA	GGCGGAGCTT	GCAGTGAGCC	GAGATCGCGC	CATGGCACTC
91381	CAGCCTGGGT	GACAGAGCGA	GACTCCGTCT	CAAAAAAAAA	AAAAAAAAA	AATTAA
91441	ATATGAAGTT	TTGAAGCAGA	AATTATTTTG	TCGTATGTTC	TTTCATAAAT.	TTTTTGCCTG
91501	CCTGCCTTCT	TCCTTTGTTA	CAGAACTCCA	ACACTTACCC	AAAGGTAGCT	GTTGGGTCAG
91561	GGTTTCTGTA	CTATAGTCCC	TTCTGTGGTG	GCCAGAAATA	TGTTACAGGA	AAGAGGTCCC
91621	CATCCAGACC	CCAAGAGAGG	GTTCTTGGAT	CCCGCGCAAG	AAAGAGTTCA	GGGTGAGTCC
91681	GCAGTGCAAA	GTAAATGCAA	GTTTACTAAG	AAAGTAAAGT	GGTGAAACGA	CAACTACTCC
91741	ATAGACGGAG	CAGGACATTC	CCGAAAGTAA	GAGGAGGAAG	GCATCCACCC	TAGGTACAAT
91801	ACTTGTATAT	ATGGGGAGAT	GTGCTCTGCT	ACAAGTTTGT	GATAAAGGAT	TAATTTTCTT
91861	AGTTACTATA	TTTTGCAAGA	ATCAACATTA	TTATCTTTAA	ACAAAATTAA	GAATGCCTTT
91921	GTTCTCCAGA	TATAGGGATA	TCTGGACACT	CCTAAGTCTG	AGTCTGTTTA	GTAAACATTA
91981	TTTATTTGTT	CCCTTAACCG	TAAACATCTA	GAAGCTAGGA	ATGACTGACT	TTCTGGGAAT
92041	GCAGCCCAGA	AAGTCTCAGC	CTCATTTTCC	TAGCCCTCAC	TCAAAATGGA	GTTACTCTCC
92101	TTCAAGTAAC	TCTGACACTT	TTCTTCTCTT	TTTTTCTTCT	THEFT	CTTTATTTT
92161	TATTTTTAT	TTTTGAAATA	AGAAATCAAG	AATACTTGAT	GTTTCATCTA	AAACAATACC
92221	CATAATTGAT	AAGCCAAAAC	AAAAACCTAG	GTCTTCTAAC	TCAAAACTAG	GATGTTTTCC
92281	TGTCTCTGCT	GATACTCGGC	TGATCGTTAA	TAGGTAATTA	ACAAACAAGC	CTTGCTATGT
92341	CCCCCTCAGT	TTATTACCAT	TAGATCATAT	GCCTACTGTC	AATCATATTA	ATCCACAACT
92401	ATGCATTTCA	CAAAACTTGC	CATAAAAATT	CACAGGTTTC	CCGCTTCCCT	CGAGTTTTCA
92461	TTTCCGAAGG	GTCCCATGTA	ATATAAAACT	TATATTAAAT	ACATTTGTAT	CCTTTTCTCT
92521	TGCTAATCTT	TTTTTTTTTTT	TTTTGAGACT	GAGCCTTGCT	CTGTCACCCA	CCCTCCACTC
92581	CAATGGCGCG	ATCTCGGCTC	ACTGCAACCT	CCGCTTCCCA	GGTTCAACCCA	ATTOTACTO
92641	CTCGCCCTCC	CGAGTAGCTG	GGACCACAGA	TACGTGCCAC	CATCCCCCCC	TA A TOTACTIC
92701	ATTTTTAGTA	GAGACAGGGT	TTCACCGTGT	TGGCCAGGAT	CATGCCCCGC	TEST
92761	GTGATCCGCC	CGCCTCGTCC	TGCCAAAGTG	CTCGGATTAC	AGACCTCACC	CACTIACCTC
92821	GACCAATCTG	TCTTTTTGTA	GAGGGGCCTC	AAGCATGAAC	TTACTCATCC	CACIGCACCC
92881	CAGAATTTTC	TTTTCCCCTA	CAATATAAAC	ATTAATTGTA	ATCTTATCAT	GIGAGAAAAA
92941	TTGGTGACCA	ATCTTACAGA	AATTTATCT	TGTGCAAGTC	TATCCAAACC	TCAGGACATT
93001	TCTTCTATAA	GTGAGATTGT	ATTTCACTTT	TCTAGTATCC	TATOCAMACC	AATATGTAAA
93061	TTCTAATGAT	TATTTTCATT	ACTGCATTTC	ATTGTAGGGA	ACTACATA AT	TOOCCOMMAN
93121	TCACTGACCT	TCGCTTTTTA	מממדדדמממ	CCATGTTACC	ACTAGATAAT	TUTTE
93181	TTCTCTACAC	ACAAGATTGC	TGTAAGGGCA	AAAATAGAGA	TACCAATCAT	TTTTCAGTAT
93241	ATATACATAT	TTTCATTTT	AATACATCTT	ACCAAGTTGC	CTCCTCALAG	GCATCCATTG
93301	CTCTCACCAA	CAGGGTGTTT	TTTCCTCACT	TCCACAAATG	CTCCTGAAGG	TCTGTTTACA
93361	TAGTCTGTTC	AAATTGCCGA	CATCAACAAT	TAAATCTCAT	TOTTOTANO	ATTIGGTTGT
93421	CAATTATTGT	TTGAGACTGC	ACATTTTCAT	AATAACATTT	CTTCTATTA	ATTTTTAAGA
93481	CTCATGATTC	TTGCCCATTT	TOTTTTCCCA	TGTTGCCTTA	CITCIATIAT	GGTTTGATTA
93541	TAGCTCCATC	TATTANANA	TTATTAACTT	TGAGGGCTTA	TCATACATTAT	TITAAATAGA
93601						
93661				GGAGTTTCAC		
93721				CTCCGCCTCC		
93781				GGCAAGCGCC		
				TTGGTCAGAC		
93841 93901				GTGCTGGGAT		
73701	CCCGGCCACA	TITCIAAATT	CITTATAAGT	ATAAATTCAT	TCAATCTTCA	CCAAAACTCA

Figure 8 (Page 29 of 73)

Erdnre 8 (Page 31 of 73)

DAADTDDTD	DAABOTBAAS) TTATDATTAT	TATTAADDDD	DTTAATDDTA	AAADADTTTT	τθεοοτ
ATDDAAAAA	TTODIADIDA	S SSTOCARAGO	AABADDAATA	AAADTDAAAD	TODDODADOA	τσεοοτ
DDAAAADDD) DTAADADTDA	DTADAAAAAD	CACTCCTTTC	OTDADDOTDD	TTAAADATTT	192001
ATDTATAAA1	TOTOOADDA/	SCCCGATTTG 1	GAGCCACTGC	TACGGGCAT	ATOOTOOL	τοσοοτ
AADDDTDDD	DIDDIEDODE	DOADTOODDA	ADDITOTED TO	DOADOTOTTD	TADOTOTITO	τυτοοτ
DDADADADDT	. DATTTTATTE	DTTTTTTG (TODDDDDDTDT	LILLCLLLL	TTTTTOOOOO	T8000T
DOAADDATDA	S SET SOUND A SET	TOAADDTOAA	TOADOTOTTO	DADTDDADDD	TOOTADODAA	100001
つかしつりついつて	. DOTOOTOOAA	A DOTDADTDAA	DIDIATOTO	DIDDIDIDAD	9TO9OACCOO	19666
DTDTDTDDDT	. DTTDTDADAD	ADADITITOTO	CCCLLLLLLL	DTDDDDDTDA	CTTCCCTCCC	T0666
TTTTDDDTTTA	TDDDDTTTA	DOTTCACTGG (AATOTTOĐAA	TATTOĐAĐAĐ	TATGAATCAA	T#866
TOOOTOTTTO) TTOOOOOTAO	ATOOSTTOAA	ACTCTCCCCT	TATADTAAAA	DOTOTILITIA	18466
TTTADDDADA	AADTTTOTTA	TODADTOTO	ADADTOTTO	TCTAAACAGA	ADDADAATTO	12766
DTDDTDADA 2	DABABBADAD	TAACTGTAT	CCAGAGAAGA	AASTASTSTA	Apppetrocor	19966
TOADTTADAA	TTDTTDATTI	CAAAAAGGAA	TTAAAAAAAA	ATTAAAADTT	TAAAAADTAA	10966
TOTOAADTOE) DITITODDDAM	DDAADTTAAA	TOADAAADDA	TOOADBBBAB	TOAASTTTOT	T \$ 566
TTASTSSSAT	TOTITIOADDD	TOTAAAAAT	DAADSTAAAS	DADTTDAAAA	ADTOTTOTOA	18766
DTTADTDDTD	ADADADOTTA	DAAADTOOTT	TACTCTCAT	TTTDTAADDT	ATTOTOTOTT	72766
TATDADTTTT	ADDDIAATAD	TAATATTT	ADAAADTDAD	TOOTOOAAOA	SERVERSE	T9E66
DADATTDADD	DTAADADDDD	TTTDABATDB	DIATOTITED	TOTTAADDAD	SADITIONAL 2	10566
DODIATOTIT	DOTOTTOADO	AAAADDDATD	TTDDTTDDTA	TTTTĐĐATTT	TOATAATOAO	T 7 Z 6 6
ATTODATODA	ADAAADDDAD	ATDDADTATT	DTDDDDDDTA	ADATTTTOTO	TOADAADAbb	18166
ATATTƏƏTƏT	TOOTTOTTTA	DOADOTOADT	DTDDADTTDT	TOTTABAADT	DIDIDITION	12166
AADDDDDDA	ASSSTDAST	DADADAATDD	TAATOTABOA	TOOOATATOT	SECRETARA	T9066
AADAADTTƏÐ	ATTODIATIT	TTADDDDTTA	AADDDDADAT	ADAAADTOAD	AAIIAAAAII	10066
TTAĐAĐĐAĐA	TTTTAADADT	ATTTATTTA	TATATADDDA	AAAbAbburba	ATTACACACA	
AAADBBTDAT	TAAAADDTAT	TTĐĐẠACCOT	CCCTTTGCCC	OTTOAAAAAA	DARJUGAGA	₹\$686 ₹8886
TOTTATAATA	ATOTOOOTAA	DABADDBABA	DDADAADTDD	TTOOATTTTA	עזכופוכוופ	12886
ADBBATDDDT	TTABAADDAD	AAAAADDAAA	ATATAADDTA	CTCTATCTC	DIDING	
TOTOTADITE	DADDTAATTA	ATAADDTDDT	TAAABASTSS	ATABATBATB	AASA I ASASA	19486
DDADADAAT	ATDADTTTDT	CTGTCATCC	TODILLITICAL	TTOTADTOAD	ATTITITION	10786 14986
ADBABBADDB	ADDDDTDDDA	CGCATCAGCC	ADATTABBBT	TOTOAAAOOO	Tooberroo	T8586
COCCTOTABL	DAADTDTTDT	DOTOAADOAT	TabtoasAcc	DOTTOTACOA	D.L.L.P.P.P.L.D	T Z S 8 6
ADAADTDATT	TTTATOTTTT	TTAATOĐADO	SSSASSASSA	2227255424	TIMESSESSES	
TOADOTATOO	DADTTDDTDD	TOTTABODAA	STTBBBASST	STATESTATE	2012A21200	T9#86
DTDTABDAD	DTDADDTDAD	DTODDADDD	TTOTOTOTE	JAJJ LAAAJ L		10186
TTTTTTTA	DADDAADATT	CAGCGTTGGC	ADTOTOTIOA	TASSTAAAST	AMOIDIDIT.	79586
DADADTDDTD	TOATTTTDAD	DADTAAAAT	SAAASAAAAA	ADDAMAND	O I I JACANON	18286
DYDABATDAD	AATADAAADD	DAADTDTTAT	AASTTSAASS	SAMAAJITIT	WINITALA	78227 78227
DAADTDDDAD	TTDTTATODD	DATTTDTADT	SATTABTBAS	SEASTAAAA	AMMULLITIO	T9T86
ATSTTSTAAS	TADTADDTDT	ADAATTTTTD	AATABASTOT	ATURUNDANA	ITOIOIMATA	10186
TDADTOTT	DODITOTIONA	TODATOTACT	TOTAATTTAT	רפזפכטיטור	AMDAJOIAIA MIDINIA ATA	T\$086
TCTTATTTT	AAAADTDDTT	ATTTADTATO	TTTTOSTOTA		ATATICANA	T8616
DATTTAATDT	TOOTAAATOT	STSAASTAST	SOSTSTACAT	AIAUUIJUIA	TAADTABTTA	12676
DDTATATAA	TOADTITTO	ADDATTDADD	PTOTOTOADA	AUAL JUAUL	5271201A12	T9846
DITITICACTG	SETTETCTEC	ADAAADDAA'I	SOCIATION SECTION	ADAMADATO	Aフラフィンシャンク	10876
TOTTATTODA	TITATOTATT	ADTITIADAT	CACITATIOAS	CASCASCAS	422121111	T \$ L L 6
CCTTTCTTTA	DOTATTOTTT	TTTOTTATOT	TTATTOTAD	CACCACACA	インフェン・・・・	T89/6
DADTAAADTT	CAGGTCTGAA	TTOOSTTOOT	TOTTOORED	OTIONIOTI	インサントサンサン	17946
AADOTATTT	ADDADDTADA	DAAADITIDAA	PLUITING	ひょうりょすむひつみ	ンンAAC+2	19546
DATECTETAC	ADATTDAADD	ATTIBILITA	DATD I DU I AM	AMJULLAM		T05/6
DATOTODDA	TOTATTTTAA		PALL J J J PAL J	* * * * * * * * * * * * * * * * * * *	PDA 477TATA	1886
TTOOTOOTOA	TOTADOTDAD	TOTTALADIO	イベルトリング	DA ADTITITIO	TCTAGT	18676
TTTTTDADAA	DAADADTADD	AMUST AMUST	ATTACTACTO	4557726	PTADDTADTD	12576
DATDATAATD	PTTTTTTTA	ADTACODA A	シャルス スキャンシナ	TTSOTTOTTO	anabandaTA	T92L6
			TACTGA A DETO	TOADDAADTT	TTTACTGTTT	31207

102607	1 CMC 1 mcm 1 C					
103681				AACTGTCAAC		
103741				TAAATTCATT		
103801				CATTTTAAAA		
103861				CTTATACAAG		
103921				TAGAGATGCC		
103981				TGAAGGTCAA		
104041				CTTTTTATAT		
104101	AAATGTGTTG	GAGGTACTGG	GTCTGACGAA	TAGCATAAAA	GAATAAAGTT	ACATTACTGT
104161	CTGAGGATCA	GATGGACAGG	GGGTGGTAGC	TCAGTCCAGC	TATTTTCCAC	TCCCTCACTT
104221	ACATTCTTTG	CCCCCTCCTC	AACAGAACAA	GGATTCTGCT	GTAACTCTTC	ATTGACAGTT
104281	GATATTTAAA	AATTAACGAA	TGGATGAAAT	TCTCATTTGT	GAAAGAAAAT	TTATTGAGCA
104341				AATATTATAT		
104401				CAAAAAGGGG		
104461				ATTCTTTTGA		
104521				TTCTTTCTGA		
104581				GAAAAGACAT		
104641				GGAGTCATAG		
104701				CTTTAATCTC		
104761				GCATTTAGCT		
104821				TAAAATTTAC		
104881				GAATTATCAG		
104941				AGATAGAGAT		
105001				TGAGCCGAAG		
105061				AAGTAACTGA		
105121				TTACCTTGTG		
105181				CTTGTGACCC		
105241				TCTACCCAAA		
105301				CAGCCCGCCT		
105361				GGTCTCTTCA		
105421				CAAAGTGATG		
105481				GTTTTGGTTT		
105541				GGGGTGCATT		
105601				TGATTTAAGA		
105661				AACTACACTC		
105721				AAAAGAGAAC		
105781				GGAATCTGTC		
105841	AGGCTGTGAT	AAAAAATTT	CAATCCTTAC	TAATACATAC	ATAGTTGCTG	CTAGCAATAT
105901	AGTGTTGGGA	GTAAAAACAC	GAAAATGAGA	GTTCAGGACA	ATATCCCAAC	TCTGAGCAGA
105961	TTTTTTTAAG	TAGTAACATC	AAATTAAA	CCATATTATG	TAATATTTAT	TTCTTTTCCA
106021	CAGTCTCTTC	TCATGCCTCG	TTCACATTAG	CTAATTAAAA	GTCCCCTGAG	TATCATCATA
106081	ACCCGATTTA	CAGATGAAGG	CACGGTTGCA	ATGAGCTATC	ACCCTCTTCT	GAATGAGACA
106141	GTACAGTGTG	AAGGATAGCA	AAACTCCACT	CCCATCCTCT	TAGGGCTCTG	GCTGGACCAG
106201	CAAATTAAAT	TAATGTAAAA	TGGATTAACA	GGAGAAAGGT	ATATGCATTT	ATTTAACACA
106261	GGTTTTACGT	GACACAGGTG	CTCTCATAAG	GTAATGAAAG	CCCAAAAAA	GCAGTTAGCT
106321	ACTTATATAA	TGAATTGGAC	AATTAGTAAA	ATGTAAAAAT	GCGCTAAAGC	AAAGGGATTT
106381	AGGCTAGAAT	ATATAACTGT	GTAGAGAAGC	GCCCAGCAAG	GGCTAGTGCA	AGGTTTGTAC
106441	AGAATTCTCT	TGGCCTCAGC	CTCCTATCCT	TGAGAAGAAT	GTTGCTTTTT	TTAAACTACA
106501	GTGAGAACAT	CTTTCATATG	AGAATTTCAC	CTACTGCTTC	TAAGAAACAG	GTCAGCTTTC
106561				ACGCCTGCTC		
106621						ATATTAGTGA
106681						CAGATTCAAA
106741						CTATTGTGTT
106801						GAGGCAGCAA
106861						CGTGTATAAA
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Figure 8 (Page 33 of 73)

Eigure 8 (Page 35 of 73)

5T5555TT56	DECASTSATA	STADBBOTTD	DTDTADDTDD	Ababbababa	DTADDDDDDT	113341
つエンエンンンシン	STAATDADAD	DABABABADT	BDDABBTTDB	TTTOODADAA	AADTOTODDT	113281
DDTDDAAAA	TTDATTTDAT	DDADATTOTT	TOOTTAAATT	STOTITIOS	TTADDTAADD	113221
DIDDIDAAAA	DOAAAATTAD	DATDADITIO	DAADTTTTDD	TTADDTAAAD	SCGACCGGCC	τοτεττ
COACTOASTS	DESACATTAA	DTTODTDAAA	DTDTDDDADT	DOSCOSOLT	AATDDTDTTT	τοτεττ
ADTOOTOAAS	DODIABLOSS	ADDESTIBLE	DDDDDTTADD	DOADADATDO	TTTTTATOTT	113041
TTTDTDDDDD	ರಾಶಾಶಾಗಾರಾ	DOTOBOBAAO	ATTAĐĐĐTĐĐ	ATOTOAOACT	TOOBAOTOOD	112981
TOOTOTITE	. DAADTTDDAD	TOTOODTOTO	DAAADTDADT	TADDTOTADD	DADATTAGAT	115351
9A99TDA9A2	DODOTOTOTO	TOTARDODAA	CLLLLLLLLLL	AADOTDAATT	TTATTAADAD	115861
つずつならずつずする	DODAAATDTA	AATTAAĐAAT	TODOTOAADO	ADTTTDDDAD	ATOBABADAA	TTSBOT
TTOTTTT ATT	TOODOTAAOO	ASTITODBAGO	AADDABBDTB	DDADTADADD	DOTDDDADDD	てサムててて
SOTTASTS AT	PADDDDDTA	AAbbADDDD	AADDTDDTDD	DDAATTTDDT	DDDDTDADAT	112681
SSSSASTSTA	CAGGGGGAA	DODODAAAAD	TOADADTOAD	ASSOSASSTS	うようううううしてつ	115621
222ATT22AT	· DTATATATT	STABSSAAAB	DATAADA DAD	TDAADDADAA	SOAASTSAAS	115261
AADDDTTDAE	TADABADDTB	DADDATTDDD	DOOOTTOOOA	AADDDATTDD	DDDDADDTT	TOSZTT
Sections	TTDDADAADD	DADTOTTOTA	DOTOCOSTAD	TOOTTOOOOO	CGCCGGATGG	175441
4842288922	SACCECCACE	ADTTOBOOTA	DADADDADAD	TADATOCTO	TOTOADDDDD	115381
COACEASTIC	DADAAAAADD	TOTTBOADTA	DECACTODDA	DADTDDATAT	อววTวอTอ วอ	TTS3ST
9TT2929TT2	TOTOAOTOAO	DDTADDTADDA	SSABATBSBS	DAATTDDDDD	TOOOTOOTO	192211
554AATADD5	AAADDDDDAA	TTDDAADDAD	TAAADADDTD	TODAAODTAA	ADADTTODTT	112201
TODODAAAA	, อ Tวววออวออ	DITDATITET	TTTOTTOD	ADATABBTAA	TTCASTTCTT	TTST4T
DEADTTADAD	ATTAATTOTA	DTTADTATDA	DTDBADTDTA	DDATTTTTDD	TTTCTTAT	115081
CONTRACTOR	- DAAADDDDAD	DTDADTDATD	TATADDDATD	DODITODITED	TATAADDTTD	ττοσττ
ADADITIOTIT	DIDADDITIT	TTTDATDADA	ATTTTTAAAD	ATAATDAAAQ	DTTCCCCTTG	196111
エンエエンシンシタン	DODTTADTTA	DTDDDDTAA	DDATDDADAD	TTODTTADTO	TTOADTADTD	106111
DATTATAAAD	DACTOCOTTO	ออ ฐวววรวออ	つつてつつつつててつ	DATTOTTODA	ATAADTDTTT	111841
	DDADAAAATD	DILLILLILL	TTƏƏTAƏAAA	ATDEDTAATA	DTTTATTTTA	τθζτττ
DD11710107	AAATTOATAD	DDADTTADTA	SATSASTSTS	DTTATDATTT	AAATA DDDDT	TZLTTT
SOTTATOTOA	AAATDADATA	ATTITITIOTOD	CAATTTTCAG	TTTDDDATAA	DTOTOADDAD	199111
ATT ATT SESTING	DTDTDATTTA	ASTOCOCOTOT	AATTTTAĐAD	AAAATOTOTO	TTTTDDDTDD	το9τττ
AAAAASTOAS	AATDDDDDDD	DAADTODITOA	ATTDDAAAAA	AAAADDDDDA	AATDDDTDTT	TFSTTT
TASSITTAGA	DODADTATOT	TTATAADDOT	TTAATTTAAD	ACCATCATCA	ADATOTTDAT	187111
T-1444415T5	DATITIAADI	AADTTĐADAT	CCACTTTAGG	CAPARTTTGC	ADDATOTAAT	177451
TTATASTTAT	TTDDADTAAT	ATATOTATOT	DTATATAAA	ATATATADDT	DTDTDTTTT	τσετττ
4つエ4つむエエン	TOOOAOOATO	STSTASTSSA	ADDATDETAT	CAACATTGCC	DADADADTTD	τοετττ
2255755424	CAGGTTTAGC	DITIODDADDD	DAAADABBTD	ADTTODDTAO	DATDESTATS	111241
AAAATTTTTT	ADAADDADTA	SOTSAATASA	DAADDTDAAT	TDATOTADAT	DDAADTDTTD	τεττττ
421251117	CACATCCCTT	TTĐẠAĐTĐƠA	ATDDADTATD	DOTITIATADA	ATDADADTAD	177777
ADTO ATTT AT	CATTTGTGAA	DDADADTAAD	ADDADAAAAA	CAATGGTACC	AATATOOTDA	τροτττ
2744457755	APTECTOR	TODAADTOAD	TADDTADADA	DAAATAAAAD	TODABBADAA	τοοτττ
TCAGGTGGAA	DTDAAAADTA	DOTTOTAAAD	AADTDOTTAA	TOTOTTOATO	DDTDDAADDD	τν6οττ
TOTOTOTI	CATTGTCAGT	DIDITITAAD	TATODAODIA	ADADDITATDA	AADTOOTTAD	TTOBST
TAMMATITOT	AGGCTCATTT	DOATADDITOD	DADDDADTAA	AATADAĐAAA	DAATDAADDA	170857
TOWNIDITE	ATAAATAADT	ADADTATADD	DTDTATAADA	ADTTTADTAA	DDDDATDTAT	T9LOTT
TO A STORTS	AAAATTAADD	DAAADDADTT	AADDTOTOAD	TTADATOTOD	DOADADADOT	τοζοττ
ATODAMANO	DETATADODT	DOATATOADO	AADATTOTTA	TTODDTDOTD	ATSBASSASA	179011
ADTADARTO	GAAAGGCCAG	TATOOTAADO	ATADAADTAT	DOTAADDADT	GTATGACATA	TBSOTT
TEANART	DTATDDDTDD	TODIES	Debaabaat	DADDATTDTD	GACTGCATGA	ττοεστ
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TAASTSTSTA	DDAAADTAAD	TTSTTTSSAS	DTATTTDATD	DADAATTTDD	AUTUDEDALA	TIOSBI
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116701		CAGGCTACGA				
116761		GCAAGGGAAC				
116821		AGAAGGCGTC				
116881		CAACGGGTGC				
116941		CTCCGAAAAA				
117001		CCAAAACTGT				
117061		CCAAGGCGGC				
117121		CCAAGAAAAA				
117181		CCACCTACGC				
117241		AACACGCCCT				
117301		ATCTAGTAAG				
117361		AGTTAAAATC				
117421		ACTCCAAGCT				
117481		AGGCTTTTTA				
117541		AGTAGCATTT				
117601		TATGGCTTAA				
117661		ACGGTGACCC				
117721	CCCCGCCAGG	TAGGCCTAGC	TCGCTTGCTT	TCTGCAGCGC	CATCATGACA	AAGCTTTGAA
117781	ACGCAAAATG	CTTTCTTTGT	GCAGCGCCTT	ACCATGGGTG	CACTTACGGG	CTGTCGACTT
117841	GGTTTAGGCC	CTTGTCAGGA	CAAAGGAGCT	TAGTTTGTTG	GAGTTTTAGA	GCTGCAACCC
117901	AAAATCCCTT	GCTCGGTTTC	TCTGTTTTTA	GAAACGGAAG	CGCCCTGATT	GGATATTTGA
117961		GCTTAACTGG				
118021	GAGCCCACAC	ATTCAAAACT	GAAGATCCTT	TTCTCAGAAC	TGCCCCTTTA	AGCTTTTGCA
118081		TGGGGGTCAG				
118141	TATATGATGA	GCCAAGTTTA	CTCACTTTTA	CTTAGTGCAG	TTCAATTCTA	AAAGTTTATT
118201		TGCATATGAG				
118261		AGCTCCTTCC				
118321		TCAAACTAAT				
118381		GTGAGAAATA				
118441	TCTATATACA	TGCATACTTG	TGGTTTTGTT	TTTTTAAAAA	AAAAAAAA	AAAACACGTT
118501	ATCTTTTGAG	ACTGGGTCTC	AGTCTGTTGC	CCAGACTGGA	CTGCAGTGGC	ATAATCACAG
118561		CCTCCAACTC				
118621		ATGCACGCAC				
118621		GTTGTCCAAG				
		TTACGGTCAC				
118741	TACCCTTATT	AAAAATTTAA	ACAAAGCCTG	GACGCAGTGG	CTCACATCTG	TAATCCCAGC
118801		GCCAGATGGG				
118861	ACTITAGGAA	ATCCCATCTT	GACAAAAAT	TTAAAAAATT	AGCAAGGCCC	AGTGGCACGC
118921	ACAIGGIGAA	CCAGCTACTT	GGGAGGCTGG	GGTGGGAAGA	TGACTGGAAC	CTGGGAGGTA
118981	CACCCTCCAC	TGAGCAGAGA	TCCTCCCACT	CCACTCAAGC	CTAGGTGACA	GAATGAGACC
119041	GAGGC I GCAG	ACAAAAATAA	TTTTAAAAAT	TTACAACGAT	GTTATATACA	CTTCTGCATG
119101	CAGTCICAAA	CTTAACCAAA		ACCCTGTCAT	GAAAAAGAA	ATCCTTCACA
119161	TIGCTITIC	TAAGTTATTC	NTCCNTTCT	TATTCATAG	CATTGATGTT	TCCAGTTACC
119221	TGGAATAGCA	ATGGTGCAAT	. MICCMITICI	TALIGATARG	AGATTGCTAG	GTTTTAGGTT
119281	ACTGCTGAAG	ATTTTATTTA	TGAATAGAAT	TCCAGGGCIG	CACTCTTACT	CTCTCACCCA
119341	GTATTTTAT	C ATTTTATTTA	A TITATITATI	CARCTTTCC	CTCCTCACTT	CAAGCGATTC
119401	TGGTGGAGT	CAGTGCCATG	ACCICAGTTC	TOTAL COLCE	CTCCTGAGIT	COCTGGCTAA
119461	TCATGCCTC	r GGTCTCCCGA	GTAGCTGGGA	TIACAGGCAC	CIGCCACCAG	GCCTGGCTAA
119521	TTTTTGTAT	r TTTAGGAGAG	ATGGGGTTTC	ACCATGTTGG	TCCCAGACTGGT	CICAMACICC
119581	TGGCCTCAA	G TGATCTGGC	ACCTCGGCC1	CCCGAAGTGC	TGGGATTACA	GGTGTGAGCC
119641	ATGGCGCCA	G ACCTGGACTT	TGTCTTCTGT	TTCATCAGTC	CTTCTGTTGG	TTCAAGCACA
119701	GTATCACAC'	r gaagactga7	GATTCTATAT	T AAATATGGTA	AAGACTGTAC	ACCCTAACTG
119761	TTCTTATTT	LATTTTAATT 1	GGCAATTTT/	A GATTCCAGCT	TTCCAAAGAA	TTGTGGAATG
119821	CTTAGAGCT	A GAGAAGCCTT	GGAAGTCAT	r tagtttttgt	TTTGTCAGAG	AAAATTCTGT

Figure 8 (Page 37 of 73)

Eigure 8 (Page 39 of 73)

DATDDADDAT	TAAAADADAD	TTTDDDTADA	TAAAAADAAD	TTAAADTDAD	DATDADADAA	126301
DEABATTAAD	ATDTADTADA	AAAAATDTTA	ATƏASASTƏT	GGCAAGAAAA	ADDITOAADAD	156241
DDADTDADDT	ADDIDAADID	TODATTTAAA	ADDIADATET	DAAAADTTDA	TAATDADATO	756787
DTTTTCATTA	TADATTATTD	ADATTADDTD	TTTATATTTT	ATADOTTATA	TTTTTTTTT	756757
DTADDTATTT	ASTABBBTTB	TDATTOOTTT	DTTATAADTA	DTTATTAATT	CACCAATTAT	120921
DTATDADTTA	DIDITOTATO	TOOTTTOADA	ADTDATDDAA	AATTTTDDAD	ADDDAADTTA	150921
TTTADDADTA	DTTDDAADAA	CAAGCCCCTG	DTTTTATA DA	TTAAASTTAA	TTTCTTTAAA	178941
TODTTAAADT	DTATTOTTT	TTOTATOTAT	TOTACTTTTT	TATĐAADADD	ADDAAADTDD	188571
DTDDADTADA	AĐAĐTĐTĀĀĐ	TOOYOUT	Sestes	TASSOCIATE	Aoroobreo	128521
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129601	GCTAATTCAG	TTTTCAATCA	TCATTAAAAT	TTTGTTCCTA	AATATATGGC	CATTATTTTC
129661	CACAACCACA	CTAAAACTTT	ATTACCTCTG	GCAAGTGACT	ATGCAAGTAA	CTAAGAGCAA
129721	AAATATCCAC	AACTACCATT	TGAGCTATCA	ATTTAGGGAA	AGTCATCTGG	CTATAATCTA
129781	AGTGACCCTC	CACTGAATGT	CAGTATCTTT	GCATATGTGA	TTTAAATCTG	GGCCTTCGCA
129841	ACACCATGAA	CTGTTCTTGT	CTTGAATATC	CAGATTGAAG	GAAATAATCT	GAGTAGTTAC
129901	GAGTCCTGAA	GCTAGAAAGA	TGGAAACCCC	ATTTGCTCAT	CAGAAAGCCT	TAGAGCTTGG
129961	GCGCTGGCGG	GTCCTGTCTC	ACCGGGACAG	AGGGGCTCTT	TCCTCCCCAT	CTGATAGTCT
130021	GATAACTAGA	GAAGCCGGCC	AACTTATTCT	CCAAGAAGGA	GCCATCTTAG	TTCCTCCTGA
130081	AATGTTCATA	TTTAGAAATT	ATTGTTTGTC	AGTAATTTAA	CCCCTTAATG	GGCTTGCCTT
130141	GTGGTCCATA	CCACTGAGTG	CAGAGCTTGC	CTGGAAGAAT	TGTGAGGGCC	ATTCCATCTT
130201	CCAGGCAGTA	GAGTTCAGTA	CTTCTTTAAA	ATTGCTGCTG	AACTCTGTAT	TTGAAAAGAA
130261	AGAATCATTT	GGGTGTGGTA	GCTCACACCT	GTAATCCTAG	CGCTTTGGGA	GGCTGAGGTG
130321				TGAGACCACC		
130381				TAAATACAAT		
130441				CTCACTGGAG		
130501				ATTCAATAGG		
130561				TGTAGACACC		
130621				GAACTTTTTC		
130681				GTTTCTGAAA		
130741				ATAAGCTAAT		
130801				GACTTCAAGA		
130861				AGAAGGAATT		
130901				CCTCGGGTTG		
130921				CAACCATCGT		
				ATATATTTT		
131041				ATTTTTATAA		
131101				TTTGCTGTTT		
131161				TTAAAACTTT		
131221				CAGGCTGGAG		
131281	TITITIGAGI	CAGAGICACA	TCCATTCAAC	CAGTTCTCCT	GCCTTAGCCT	CCTGAGCAGC
131341	TTACTGCAAC	CCTCTCC	1 GGATTCAAG	GCTAATTTTT	GTATTTTAG	TAAAGACGGG
131401				ACTCCTGACC		
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131521				GAAAGAATTC		
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131641	AATTAATTT	ACAATTAATA	IGATTIIGAA	ATTGGGTATT	TATTACCATA	THE STATE OF THE
131701				ATTTTCATCA		
131761	TAGAATTTCA	TAATTTATAA	AGCTACAAAC	TGTATGTGAT	TCANCATTCC	ACTITATETC .
131821	ATAACTTTAT	GCAGTTACAA	GTAGAAATAA	AATGTTCCCC	CCTCTTTTT	CCCCCAAAAT
131881	ATTATAAACA	AGTGTAAAAA	ACAAAATCAC	TAAAACACTC	A COTOTITITI	TOCOTACATT
131941				TAATCAGCAG		
132001	TGTAGACTAA	ATATTAAAAG	TCCCAAAGCA	AATGCATTTT	CCCCCACCCT	CCARCCACT
132061	TTTTTTTTT	TTCTTTTTCT	GAGACGGAGT	CTTGCTCTGT	CGCCCAGGCI	TCCTCCCTCA
132121	GGCACAATCT	CGGCTCACTG	CAACCTCCGC	CTCCCGGATT	CACGCCATTC	TCCTGCCTCA
132181	ACCTCCCGAG	TAGCTGGGAC	CACAGGCGCC	CGCCACCACG	CCCAGCTAAT	TITITIGIATI
132241	TTTAGTAGAC	ACAGGGTTTC	ACCGTGTTAG	CCGGGATGGT	CTCGATCTCC	TGACCTCATG
132301	ATCTGCCCAC	CTCAGCCTCC	CAAAGTGCTA	GGATCACAGG	CATGAGCCAC	
132361	CTACTGACT1	TTATCCAAAG	AAAATATAAG	AGCTCTTCAT	CATAACGTAT	GITICIIGCI
132421	CTTGTTATTA	A AATATGACAC	ATTTAGACTT	AAACTGATTT	GAAGGTTTAT	DACATIGITI
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132601	CTACTAAAA	AAACGAAACC	CTTCCAGGTG	TTAAGGCAAA	ACTITCCTCC	CTCTTTCTTC
132661	TATAAATCTO	ATTCCATGTT	AGTGAAATTI	CTACTGATGG	CTTTGGTTTC	CICIAIAGIA
132721	GAATAGAGAT	r cctatggcaa	AAGTCATGTC	TGACATGGTA	GCAAATAGAA	ATGGGGAAAA
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Figure 8 (Pag 41 of 73)

Figure 8 (Page 43 of 73)

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142561	ACGAAAATAA	AAATTAAAA	AAATTTTAAA	AAAAAGAAAC	AAAAGCTCTC	TARTCROCK
142621	GICCTACACG	ATAGTGAATA	AATTTTTTTG	TGTGGTCCCT	AAAATTGAGT	TCNTCCCTTT
142681	TCTGAAGTAA	TAGACGCCCA	GAGAAGGGAT	CGACTTACCC	ATCATCCCAC	ACAGGCCTTT
142741	TGGCCCCAGA	ATTCTTTAGC	AGACCGTGTA	TATGAACGTC	CTTTGCAATC	AGAGATTAAT
142801	ACTGGGAAAA	CCTCATTTAG	TATGTTACAT	GCCTAGCGTT	TTGTGCCTCA	ATATAAATTA
142861	AGAACCAGGG	ACTATTGCCC	CAATATTATA	TTTCAGGAAA	GGNACCCCCA	ACACCTTACA
142921	GTCACTGGTC	CACTTTCACC	CAGTTGGTAA	ATGAAACCAG	A A TOTAL CO	GACAAATGGT
142981	AAAGGTGAAA	ACGTTTCTTT	TATAATTTCA	CATACAACCAG	TELLEGE	TGTACCACAG
143041	CACATTAAAG	CAAGTGCTCA	GGAGTGACAT	CALACAMICI	TRATGGACC	CAGTGTCCAA
143101	GAGTTTAGGT	CTTGGAGAAA	AGAGACCCAA	GCDCDCDCDA	CACARAGECC	TGTCCTCAGG
143161	AGCGCTGAAG	ACTGAGGACC	CTGCCTGTGG	ACTGAAGTGA	CCATCCCCA	AAAGAGAAGG
143221	CGGAATATGA	CAGTTTGGAG	GGGCCTGAAG	GACTCTTCTA	TTCTCTTATCA	ACCCGATGCC
143281	ATTACTCTCC	TAACCAGAAA	AGGTATTTCA	ATTTATATTT	TOTOTALCA	GAAAAACAGA
143341	GTGATAATTT	AATGTGTTTT	AAAAAATGTA	TCACACTCAT	CCCCTCCTCT	CACTTTTCTG
143401	ATAAAATTT	AAGAATTAAA	AAATATAAA	ATCTTTTN TO	TACACATTAC	GAAATAAATA
143461	GATAACTGTG	AATTATAATT	AGTAATTAAA	TTCDAATACT	CATTATTTTT	GAGTTACAAG
143521	AATTATTTAA	TAAAACCTAT	TTAACATTTA	ATATOMATACI	GATTATTTTC	ATTITITATIT
143581	ATATTTATTA	TTATAAATTA	TTTTACATILA	ATATTTATCA	GTAATTAAAT	CTAATTGTTA
143641	CTCAAGCCTG	TAATCCCAAC	ACTTTGGGAG	CCTAACCTCC	GTAGAAGCGA	GGCATGGTGG
143701	AGTTCAAGAC	CAGCCTGGGC	AACATGGAGA	A A C C CT CT CT CT	GAGGATTGCT	TGAGCCCAGT
143761	TGTGTGGTGG	TGCGTGCCTG	TAGTCCCAGC	CATTCTCCAC	CARTACAAAA	AAATGAGCCA
143821	TGAGCCTAGG	CAGTCAAGGC	TGCAGTGAGC	CCTCATCTTC	GCIGAGGIGG	GAGGATGACT
143881	CAACAGAGCA	AGACCCTGTG	TCDDTDTDCD	TATCCACAAA	CCACTGCACT	CCAGTCTGGG
143941	CATACTACTG	ATACAGAATT	CACTACACAT	CCARACCTAC	CTTAAAATTT	AAAATGAAAG
144001	AGATAAAAAG	GAGAGTGGAA	GAGIAGAGAI	CATCAAGCIAG	TCCTATAACC	AGAACAATAA
144061	ATATCCTCTA	GCAGAACAAA	ACARCARAR	CATGAATTTC	ATGATAAATG	GCAATTGCAA
144121	GGCCAAGGAG	GGAGGATTGT	TTCACCCCAC	1GIAGATAAA	ACATATCCAA	CCCTTTGGAA
144181	AGACCCTGTA	TCTAAAAACC	AAGAAAGAAA	AAGTTGGAGA	CCAGCCTGGG	CAACATAGTG
144241	TTGAAAGCCA	TCTAAAAAGG	AMGAAAGAAA	AAAAAAAAA	GGATGATAAA	GTAGACAATA
144301	CAAAAATGAA	TAGATATTAG	TTCCCTCAAA	AATTTGATCA	GTAATTTTCT	TCCAACAGTG
144361	ACTATCTAAT	TAGATATTAG	CTACTAAAA	TAAAAATCAA	ATATCCAACA	AAAAATATTG
144421	TTTDDDDDD	AGTATCTAAG	TTTATAAATT	TGGCCAGTTA	TAAAATGTCT	TAAATTTTTA
144481	GAAGATTTTG	GAAAACCATA	ATCACAAAA	GAGGTGATAA	AGAGAAATTA	TTTCAGTTAT
144541	TTABGTTACC	TTAGAAAACT	TARAGRARA	AACTATTTT	TGTTTTCAAA	AAGTGAAAGA
144601	TRATCCCACT	AAACAGTTGC	TAAAGAATAC	CAGATGGCTG	AGCGTGGTGA	CTTATGCCTG
144661	CACCCCCCC	ACTTTGGAAG	GCCAAGGCAG	GAGGATCATT	TTAGGCCTGG	AGTTCGAGAC
144721	ATACARGAGG	ACTGTAGCAA	GACCCGTCTC	TATTAAAAAA	AAAAAAAA	AAAAAAAAGA
144781	TTATTTACCT	TTGCTAACAA	TAGCAAAGAT	CAATTAATTC	AAAATTTGAA	AAACTGTAAT
144841	TIMITIMOCI	TTAGAGTACT	CICGIGATAT	GAGATTGCCA	AATTAATACT	TTGGGTGCAT
144901	CAGGTAATCT	AAAGGACTTG	CAAATTTACA	AAGAAGTGTT	GAAGAAAAGC	CACACATTGG
144961	TACTTANA	TTGCAAAAGA	CAGATCTGAT	GAAGAACAAT	ATTTTTAGAA	TATACAAAGA
145021	TOTOTTONOO	CTCAACAGTA	AGAAAATAAC	CTGATTTAAA	GCAGGCCAAT	GACCTGAACA
145081	TOTOTICACO	AAAGAAGATA	CACAGATGCA	AGTATGCATA	TGAAAAGATG	CTTGACATCA
145141	CARARTINGG	GAACTGCAAA	TTAAAACAAG	TAGATACCAC	TGCATACCTA	GTAGAATGAC
145201	TCATTACTCC	AACACTGTCA	GCACCAAAGG	TTGCAAAGAT	ATGTAGCAAT	AGTAACTTGT
145261	ACARAGE	TGAGAATGCA	AAATGTGCAA	TCACTTTGGA	AGACAGTTTG	GTGGTTTCTT
145321	ACAAAAGIAA	CCATACTTTT	ACCATAAGAT	TCACCAATCA	CACTCCTTAG	TATTTATCCA
145381	AAGGAATIGA	AAACTTATCT	CCACACAAAA	ACCTGCACAT	AGATGTTTAT	AGCAGCTTTA
145441	COTACTO	ATCCAAAACT	IGGAAACAAG	ATGTCTTTCA	GTAGGTAAGT	GGATAACTGT
145441	GCCCAACTCTG	AATAATGGAA	TGTTATTTAG	AGTTAAAAAG	AAATGCATTC	ACTTTGGGAG
145561	AACCCCAACC	GTGGATTGCT	TGAGGCCAGG	AGTTTGAGAC	CAGCCTGGTC	AACATGGGAA
145621	CATATOROLO	AGCCGGGCAT	AGTGGCGTGA	GCCTGTAATC	CCAGCTACTC	GGGAGGCTGA
	TCACCOTCC	TCGTTTGAAC	CTGGGAGATG	GAGGTTGCAG	TGAGCCAGTG	CCACTGCACT
145681	CAGCCTGGG	CAACAGAGCA	AGACTCCTCT	GTCTCAAAAA	AAAAAAAAA	AAGAAAGAAA
145741	AGAAAAAAGA	AAAAGAAAA	GAAAAGAAAC	GATCAAGCCA	TGAAAACACA	TGAAGGAAAC

Figure 8 (Page 45 of 73)

Figure 8 (Page 47 of 73)

ATGATATCAT	ATAATDATDT	ADAADAATDT	CTTCAGCTCT	atoota400a	AACAGACAA	TZZZST
				DOTITIODDD		191251
				ASSESSET		TOTEST
				SCCCTGAGCC		122041
				DAAAADTDAD		186151
				ATTTCTTTG		126151
				DABBABTADB		198151
				DOTTATTAT		TOBTST
				STCCAGGGTG		T&4TST
				STSAAASATS		T89TST
				TODADDAATD		129151
				DDATDDDTDD		TOSTST
				SSASTOTTTS		TOSTST
				AAOTTOTOTO		てかかてらて
				CTGGATTACA		
		CTGACCTGAA		DOTODDACOD		186151
				CACCACCACA		TZETST
,				TTABBTSSTS		192151
						102151
				Teteste		THTTST
				TTTTAATAAA		TROTST
		DADTDADADO		TODADTADTO		TZOTST
				DOTOTTABTO		196051
				TOODAADTOD		106051
		•		AAATDDTDAA		T \$805 T
		TACTTO		ADDIAADADD		TBLOST
		DATDETEAAT		TTDAADDDAA		TZLOST
		TAAAAADADD		DOCTORACAT		T990ST
				AADDAADADA		το905τ
				DDDADDDDT		TFSOST
				TTATTATATT		T8705T
				DATDAATDTƏ		72005T
				TDAATODTTD		τ9εοςτ
				AOTAOTAĐOA		TOEOST
				DDADAADTDD		720547
				AAADDADDT		τετοςτ
				TAAATTTTƏT		TZTOST
	TTATTDTAAD			DAADAAATTA		T900ST
	TODBATDDTO			DTTTDTDDDT		TOOOST
				TADTTTTDDD.		146641
				AAADDATAAA		T8867T
				ATDAATAADD		149821
DAADDDTDAA	DAATAATADT	AASTOTOTTO	TODTTTATOT	DDAATTDAA D	TAAAADTTTT	T9L67T
TOTOOTTTAA	TDDTATDDDD	TOADTDATTA	DTDTDJTDAT	DTTDATTTTD	TATTTƏTTƏT	101671
AATTADTADT	DDTATTTTTO	ADTDTTTTAA	DAATTTADAA	TATTAAATOT	DAADTETATT	T\$96\$T
TTTAADTADD	DDTDDDTADD	ADDBADTDDD	DADATTADDD	TOTTDATADA	CCTCAGCCTC	185671
DTCCACCTG	TOAACTCCAAGT	TOOTOAADTT	DYDDTDDDAD	SEATTOTASS	CAGGGTTTCA	149521
ADADATDATT	TTTATDTTTT	TAATOOOOOO	DTACCACCATG	DDDDDDADAT	TDDDTJDDATD	194611
ADDADTDDDD	TTOODTOOTO	TTAASSAAST	DDDADDTDDT	DDADDTDATT	DODITION	109691
ASGCACAGCA	DITICCTIG	TTTTTAATTA	DAAATDADTA	TAAADAADTT	DADDATDTTD	146341
TADTDDADAD	TOTTATOOTO	TTTADTTDDT	ACACAACTGT	DATOTADADT	SCCATGCCTC	149281
ADDDDDADDT	ADADADDAT	TOOOTOTATA	TTTTDTDAAD	TTTDATAADD	DOTADAADDD	149221
DTOODADTOT	TDAADATTAA	TADTADTATO	TOOBAOOOAA	DATTDATDAT	TOADTTODDA	191671
TODITITATA	CAAAGATGGC	OUTATIOTIO	ATAAAADTDA	ADTTOTOTTA	TTTTTTCTCA	101671
TTATDTAATT	TTGTTTGTT	TOTOOADAOO	DDADADDDAT	AAADĐĀTTT	TODTOUTATA	140641
						507 6

155531	0000000000					
155521 155581	GGTGGCAAAG	GGAGACCCTG	TCTCAAAAAA	AAATTAAAA	ATTAGCCAGG	TATGGTGGCC
155641	CTCCACCCA	GICCCAGCAA	CTGGGGAGGC	TGAGGTGAGA	AGATCACTTT	AGCTCAGGTG
155701	GIGGAGCCAT	GATCGCACCA	CTGTACCACT	CGGCTTGGGC	AACAGAGTGA	GAGCCTGTCT
	CGAAAAAACA	AATATATACA	CACAGTAATC	AATATATATA	TTATATGTAC	CAATCAATGC
155761	TTCACTTTTA	TATATAATAT	AGATTACATC	TTATTAGATA	TATAGTATTC	CTTCTCCATA
155821	GATAGATAGA	TACAGATATA	GACATAGTAT	CCTCTATCCA	TATTAGAGAG	ACCATA CTAT
155881	ATATATCTAT	AGCATATAGA	GATGCTGTCT	CAAAAAAATT	TAAACATCAG	CCAGATCTCC
155941	IGGCCCATGC	CTGTAGTCCC	AGCTACTGGG	GAGGCTGAAA	TGAGAGGATT	GCCATTCATC
156001	CICICATIGG	TIGAGCCATA	ATCGCACTAC	TGCACCACTC	AGCCTGGGAG	ACAGAGGGAG
156061	ACCTGAGGTG	GAAGGATATA	GATATAGATA	TATAAATAA	TATGTATAGA	GAGAATATAA
156121	TATATGTGTG	TATGTGTATA	TATATATATT	ATGAAGACAC	TGGGAGAGAA	TACTATATAT
156181	ATATGTGTGT	GTGTATATAT	ATATTATGAA	GACACTGGTG	GGATGGTTTC	ATTACCAATT
156241	GGACCAAGAG	TCCAGGTATG	GAGCCAACAT	GCAATGTTGT	TGTTGACTGA	GCTGGCAGAG
156301	CACTGGTCAT	AGTTACGGGA	AAAGAAGGTC	TCCAATGAGA	CATACTTAAC	DTATATAGA
156361	AACTTGCCAT	ATACGTGGAG	AGTTCTGGTG	TGTATATAGC	CTTCTCTCAC	CAACCTAGCA
156421	ATTGTCTTCA	TCATCATTAT	AATGCTATCA	GAGCAAAGAT	GACAGCTAAA	July July The Care
156481	CCTTTCTTCT	TCTTTCTCTT	CCTTCCCCTC	CCCCACCTCT	TTCTCTTCCT	CCTCCTCCTT
156541	CATCTCTCTT	CTTTTTTTTT	TTGAGATGGA	GTCTTACTCT	GTCGCTCAAG	CTGGAGTGCA
156601	GTGGCACAAT	CTCAGCTCAC	TGCAACCTCT	GCCTTCTGGG	TTCAAGCAAT	TCTGCCTAAG
156661	CCTCCAGAGT	AGCTAGGACT	GCAAGTGCAC	ACCACCACAC	CTGGCTAATT	TATELY TOTAL
156721	TAGTAGAGAT	AGGGTTTCAC	AATGCTGGCC	AGGCTGGTCT	CAAACTCCTG	CCCTCAAGTG
156781	ATCCTCCTGC	CTCGGCCTCC	CAATGTGCTG	GGATTACAGG	CGTAAGCCAC	TGTACCCGGC
156841	CTCCTCCTTT	AATAGACAGG	GTCTAGCTCT	GTTGCCCAGG	CTGGGTACAG	TGGCGTGATC
156901	ATAGCTTACT	GCAGCCTCGA	ACTCCTGGGC	TCAGGAGATC	CTCCTGCCCT	AGTCTCCCCA
156961	GTAGCTGGAA	CTACAGGCAT	AGCACACGGG	GCTAATAAAA	TTAATTAGGT	GATAAAATTC
157021	ACTGCCCACT	GATGACTAAG	CTCTTTGGAC	ATAAAAGACA	CAGACCTTGA	AGGAAAATGT
157081	GTCTACTTAA	TTTTGAAACC	CTATTTATCA	AAAAACAGGA	TGAAAATGCA	AAATGCCATC
157141	CACATGCCAG	AAGATATCAG	CTATAATAAG	TTCCCATAAA	TCAATAAGGA	AAAGAACCCA
157201	ATAAAAATTA	TTAAACCACA	GTAAATCATG	GGTAAATCAC	AGAGGCCTGA	AGGGCTAATG
157261	GACATACAAA	AAGAATCTCA	ATCTCACTAG	TGAAATCAGA	AAAGCACAAA	TTAAGTACAC
157321	AATTAGGTAC	CATTITAAAT	CTGTAAGACT	GTCAAAATCA	TAAATTATAT	AAGTAAAGAC
157381	TCAGGGAGTT	TTGGAGGAGT	GAGAGCTCTT	ATATTGCTTG	TGGGGTAGAA	TTGGAACAAT
157441	TTCAAGATCT	GTAGTATCTG	GTAAAATTAT	GATATGCATC	CCTCACACCA	GCATGTCACT
157501	CCAAGGTATC	TCCCTGGAGG	GAACATTTAC	GGGACACAAG	GAAGCATGGA	TAAGAATGTT
157561	CACAGTAGTA	TTGTCTGCAA	CAGCAACAAC	AACAAAAAA	CCCAACTACA	CACAACTTCA
157621	ATGCCCAGTC	CACAAGGCAA	TGGATTAAAT	AAACTTCAGG	CCGGAGATGG	TGGTTCATGC
157681	CTGTAATCCC	AACACTTTAG	AAGGCCGAGG	CGAGAGGACT	GCTTGAGCCC	AGGAGTTCAA
157741	GACCAGCCTG	AACAAAATAA	AGAGATAGTG	TTTCTACAAA	AATTTTTAA	AAAATTAGCC
157801	AGACGTGGCA	GTGCTTGCCT	GTGGTCCCAG	CTACTGGGGA	AGCTGACGTG	GGAGGATTGC
157861	TTAAGCCCAG	GAATTTAAGG	CTGCAGGGAG	CCATGATGGG	GCCATTGCAC	TCCAGCCTGG
157921	GTGACAGAGT	GAGACCCTGT	CTAAAAGAGA	TAAGTAAATA	ACAACTTTGC	ATTTTCTGCC
157981	ACATTGCAAA	ATGGTGAGAG	AGTGGTTTCT	AGACTCTAGA	CTCTTTCTAT	GACTACCTTC
158041	TAGTTATGAG	ATCCTACAAC	ACTCACCTAA	CCTCTCTGTG	TCATATTTCC	TCCTCTATAA
158101	AGCAAAAATG	CCCCATATAG	AGAGGACTGT	GATATAAAAC	AAGAACCAAG	AAAAGTAAAG
158161	CTTTTCTAAT	CTGTCACAGA	CTAAAGAGTG	CTCAGTATAT	GTGAGTCATT	ATTCCTGGTG
158221	CTGGTAGGAG	TGTATGTTAC	AACTTTGAGT	CAAGTAATAT	GGTACCATAT	ATTAAGATTA
158281	ACAACAACCT	CGGCAATCCC	AGTTTGGGGT	ATGTTCCCAA	AAGAAATGAA	AGCACCAGGA
158341	TATAAGGATG	CATGGACTAG	AAAGTTATTG	TAGCAACATT	GTAATAACTA	AGTTCTAAAA
158401	ACAGCCTGAA	GCTCCATCAG	TAGGGATATG	GTTACATATA	TTTATTATTT	TCTTATGGAA
158461	TATTAGACAT	AAAAAGTAAC	GAGTAACATA	GAAGAGACAG	TGTATATATG	TTACGTTTGT
158521	ACAAACTTAG					
158581	GAAAAACCTT	GAACTTTCTC	CTTATATCCT	TTATATTGTT	TGACTGATTA .	AAATGTATTT
158641	GTTGCATCTG	CTTGAAGGCA	ATGTAAAATA	AAATAAACAT	ACATTTAAAA	ATAAAATAA
158701	AATTTATTCC	TATCACTTTT	GTAATAAAGC	TGGGCACAGT	GACTAACACT	TGTAATCCTA

Figure 8 (Page 49 of 73)

Eigure 8 (Page 51 of 73)

		ASTODIOTAC				TBTS9T
		AATAAAATƏƏ				165121
		DDTTDATAA D				190591
		DADADATODT				τοος9τ
		DTTDTAADDD				T\$6\$9T
		DTTTAAAA AD				T8879T
		DTDTADATDD				T 28 7 9 T
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		COTITITIES				101991
		TOTTTOOTAA				T \$ 9 \$ 9 T
		DDTTADAA AD				T85 19T
		DDTDTTAATD				T25#9T
ADDDABADTT	TDAAATTTAT	DOADTTTTAT	TTDDACCAGGTT	DOTODOADT	TDADTAATDD	194491
DADDDTDAAA	ATOOTOTOAO	DTDDTADADD	DADADTTDAA	ATOTOCATOT	OTTTOOTTOO	100091
AAADADTTƏT	DAATDTTTTD	TOOTADTDAD	DIDIAADITI	AADTATDATO	ADATODADTA	T 7 E 7 9 T
DITTOBADTAD	TTTTTTATA	TDAAATAAAT	AAATAAATAA	AAATTTAATA	ATDATAAAA	187791
ADTOTADOTO	ADAADDADDT	DADATDADDA	DODITABABITO	DADTDADDTT	ADADDTDDAD	164221
		ADDDADTDDD				T9T19T
		AADATAAAAT				101791
DTADDABAAD	TADADDADTD	DADDADTAAD	TODOTODAOT	DDDADDDTTT	DADDADDDTA	100091
		DDDDDDATA				186591
		TDATOTTTDA				126891
		DAAAATADDT				198691
		SCITTITGS				163801
TATATTAAƏT	DATODAAODT	DATTTTA DDD	TOTTODADTA	ATDDTTAAAA	DADADAAADT	T \$ L E 9 T
DOTDABBABA	DATTDDADAT	DDDADTADAA	ADAAADADDT	ADAADDADAA	TCAGGAGCTC	Τ 89ε9τ
DATDTDADAA	TDAADDDDDT	AAA DATOOTA	DAADDDDTTA	DDDDTTDTDD	TTACAATCAA	129591
DATTDTATAT	ADTTTTĐĐĐĐ	ADATATOTTO	TOAATTTODA	DOTOBADADA	DDATDTDTTT	195691
TTAADDTDDA	DTABBATTDA	DTTTDDDDA	ADAADATATA	DTATDADTAD	ACAGTGTTT	TOSEST
TADTTAAATA	AAAATAAAAA	ATTAAAAATA	AAAAAATT DT	ATTOATOTOT	CCARATARAC	10001
DESTIDATA	TTOTOADOTT	ADDITITIT	ATOTAAOTTT	AADDDDADTT	ATDTDTDADD	185591
DTDADDTAAT	TAAATOTETO	DTDATTADDT	TTAADDTTDA	AATDTADADD	ADTOTADTOT	163321
DAAA DTDAAD	TAAADADADT	DADAATDTDT	ATOOTTAOOA	ATTOTADTOD	DATDDDATDA	193591
TOTTOTOATA	AABABTDAAT	TOOTATTATT	TADSSADAST	GCTTTGAGTG	GGCAGGGGA	163201
TTOTTAATAO	DTDDDADAAT	DDTDDDADDA	ADAĐAATTTT	CCAGACTGGA	DADT'ADAADT	101191
AATATTDƏAA	CTTGAAGTAA	TTTTAƏDƏAƏ	DTTDADTDTT	TADATTDAAD	AATTTAĐTAT	180891
TOATTTTTDD	ADAAAAATAD	CAGTTACAAG	TATTTDADAA	ADTTĐAADDĐ	TGAAACTAGG	163021
DTTAATTTAA	TADATODTDA	DDATTDDDTD	CAGGAGAAAG	AGCCAAGACA	CTTCTTC	196791
TTTTAATTAA	DDDADTDATA	TTOOTTABAO	TOOOTTOOAD	AADDTDTADA	STSASSSASS	106791
DADDDTDTDT	AOTOADAOOT	DOTOOTTTDA	CONTRACT	AADDTDDAAD	TOADTOADTA	162841
DTDTTTTATT	ADABBBTDAD	CACAATCAGC	CACACACACA	CACACAAACA	TTTTATATAT	187231
ATTOTOAATT	DOTAAATTDD	TOAAAATTTDA	TGAACTATAT	TTAATT	SASSAAAATTO	127231
TAADAATƏTA	ADAADADDTT	DOTADTDDTA	ADDDTADADD	TTTGCAGTTT	ADTOAAAUTA	199791
DAAADTADAA	DATTTDADTT	TOADTDATDD	DIAATITDIT	TGGACAGTTA	AAAAbaaaaa	109291
AADDDTTDTD	DAADATTDDT	DATAAAATDA	AGAGACAGAA	TADTTDAATT	DATDA DDTOT	162541
ADDDAADATA	TTJƏTTTJAÐ	AATDTTATAA	CAAAAGGGCA	AAGCCAGAAA	TAAAASSAAS	185481
TTADADAADT	TODOAADA	ADATDATDDD	GTCTGACACA	AAADTAADAA	DAAATAberra	762421
ADTADDTATO	TADDTDTAAA	DOAAATDAAD	TCAAAAGGTG	DOTATOAAAD	AASTSAberra	195391
STSTTSSATS	TOTAASTOTO	TOTTTTTTAD	AGTCAGACCG	AATTOTTADA	DOTOACACTGG	105291
DATDTADTDD	TTOTOTTOAD	AATADADAD	TODDADOTOT	TCTTCTTCT	TOTOTOAGA	762241
TOATOTOAOT	DODITE ATACT	CAGACCAGCC	DODDADTTOT	DOTTADTTDA	STICLLIG	181791
TADDADOTTD	ADDIDDITIT	TADADTDDD	ADTOBBBTOT	TGCAGCAATT	CAGCCTCACC	161131
ADATBAADDT	DATDDDTTTT	TTOOATAOAA	ATAGCACTAT	TTCTCCTTTA	T.T.C.C.T.TACCA	162231
DOTADDOTTD	TATTTOATTA	DAAADAAAAT	DTADATETAE	TOADBOTADT	SOSPECERGE	190091

169481	ATGGGGTTTC	ACCATGTTGG	TTGGCTCGAT	CTCTTGACCT	TGTGATCCAC	CCGCCTCAGC
168541	CTCCCAAAGT	GCCAGGATTA	CAGGCATGAG	CCACCGTGCC	CAGCCTCTTT	TTCTTTTCTT
168601	ATAAGACAAG	TTCTCGCTCT	CTTGCCCAGG	CTGTAGTGGA	GGGCAGTGGC	ATGACCACAG
168661	CTCACTGCAG	CCTCGACCTC	CTGGGTTTAA	GCAATCCTCC	TGCCTCACCC	TGGCAGAGTG
168721	GCTGGGACTA	CAGGTATGTG	CCACCATGTC	CAGCTAAAGT	CTTCTCTCCA	GAAAGAAGAA
168781	ATGCATTGGA	ATTTAGAGGA	TACACAAACA	TCTAGCTGTA	TAGCTAATAC	AGTAGCCACT
168841	ATCATGAGTA	GGAATTTAAA	TTTAACTTAA	TAAAAATTAA	AATGAAAAA	TTCAGTTTTT
168901	CTGTTCCAGT	TGCCACATTT	TGATTGCTTA	ATAGTTGCAT	GTGACTAGTG	GCTACATAAC
168961	AGCCTCAATA	TACAACATTC	TGTTATCACA	GAAAGTTACC	TTGGACCAAG	TGCTGGGAGA
169021	AGCAATGCAG	GCTTCCTCAC	AAAAGCTGTA	AAAGAGAGAA	CTCAGGGAGT	GTGAAACTCT
169081	TTCCTATTCT	AGTTAACTTC	AAGAATAATT	GTTACCAGGC	CAGCACGGTG	GCTCACGCCT
169141	GTAATCCTAG	CACTTTGGGA	AGCCGAGGCG	GGCAGATCAC	CTGAGGTCAG	GAGTTTGAGA
169201	CCAGCCTGAC	CAACATGGCA	AAACCTCATC	TCTACTAAAA	ATACAAAAAG	TTAGCTAGAT
169261	GTGGTGGTGC	ACACCTGTAA	TCCCAGCTGC	TCAGGAGGCT	GAGGAAGGAG	AATGACTTGA
169321	GCTCCGGAGG	GGGAGGTTGC	AGTGAGCCCA	GATTACACCA	CTGCACTCCA	GCCTGGGTGA
169381					TGGTACCAGA	
169441	GTAATTAGTA	GTAACACTTA	TGCAATTGGG	TGATCTGTGA	CAGATTCCAT	TCDACCACTA
169501	TGGGGAGCTT	CACCCCAATA	TATGACTCCC	TGGTATAATG	AGTATTTTGA	ATTANACCO
169561					CTACATAAAG	
169621					TATCTCATTT	
169681	AAAAGAGGAC	TAAGAATGTA	ACCAGACCTA	ATCAGACACT	TTCACAAAAT	AATCTCTCTC
169741					TTAAACTCTG	
169801					CATTTACTGC	
169861					TAAATATGTA	
169921					TATACATACA	
169981					AATAAGGCTA	
170041					GCCCATGTAC	
170101					TCGATTTTTC	
170161					CATGACAATA	
170221					AACACTGGTT	
170281					TTCAGGAGCA	
170341					TGCAGAGAGG	
170401					CCAGGAGGGAT	
170461						
170521					TTTTGTGCCC	
170521					GGTTTGCTCC	
					TTACTTTTAC	
170641					TGCAACATAA	
170701 170761					GCCAGGGATA	
					TAGCTGGGCT	
170821					CTGGGAGCTC	
170881					AAAAACAGTA	
170941					TCTCTTGAAG	
171001					CCAGTGATAC	
171061					CCCTCCTTCC	
171121					TCTGAGATTT	
171181					ACAAAGAAAG	
171241					CCCACCCCAC	
171301					CTTTTCCGTA	
171361					TGCCACTTCC	
171421					GTGGGAGGCG	
171481					ACTCCGCCTC	
171541					AATAAAAAGA	
171601					ATCTAAAAAC	
171661	CAGAGGTTCA	GTTCACAGAC	TCTGATTTGA	GATCTTTCTA	CTTTTGCCAC	CAACTCCCTT

Figure 8 (Page 53 of 73)

Figure 8 (Page 55 of 73)

					CCAGACCCCA	178141
				DTDADTDDDT		TBOBLT
DDAATTT	AAAAATAAAD	TOAATAATAT	ATAAAAATAA	ATAAAADADA	AAAADTDAAT	TZOBLT
				TOTTTADTOO		T96 LLT
ATTOĐĐADOĐ	TOTITIODITI	ADAADTƏTAT	DOABADDETE	CACTTGTTTT	AA DTTADDTA	T06LLT
AAATDDATDD	AAATTDADAT	AATAATDATD	DADDAADADD	CCGGATAGCC	DOTITITAATA	T \$8 L L T
				DTTDDTDADT		TBLLLT
STSTTSTSS	TTTDADADAT	TTTDTAAADT	TOTOTAAATA	ADADTDODTT	TCACTCAAAT	TZLLLT
DDTADDDDT	ADTOTOTOD	DADTODTODA	DDADTOTTOT	TOOODADITO	DADDAADDDT	199441
ADDIDDAADD	DAAATATDTA	AAATOOOTOO	DTADDDTA	DTTADTTTAT	TTCTTAAACG	T0944T
TADATETEAD	DAAADDABDT	つてつてつつうつつ	TOTTOACOTT	つもてつもつつてつつ	CCACCCCACT	TVSLLT
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SSEARSTSTS	TOTTATODAA	DATTTTADAA	TCTGATAGEA	DAATTTTDDA	DADAADTADA	T984LT
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				TTOADATAAD		TBZLLT
DDDDDAAATD	TTAAAATAAA	AADAAAATAT	AGGGCAAAGG	ADDTTDAAAA	DDADTTTTTA	181441
DTTDTDTTTT	TTTDDDTDTA	TATCTTATGA	TTTDDDAAAT	DTTAAATTAA	TOADADADDA	TZTLLT
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TTOTOTOO	TTCTTTCTC	STTAADAADA	CCCACCAGCG	DODIADDATA	AAAATAJATO	194941
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181441	GCGGCCAAGG	TTCAATCCTG	GCTTAGGGAA	TGAGTACTTT	CTGATTGATA	TCTGTGTGAC
181501	CITTACCATT	TGTTGATTCT	GTTCTCTTCC	CCTCCACACA	CTGTCTTGAG	TTTTCCTCTC
181561	TCTGAGAACC	TGGGAGATTA	TCTTTGGTAA	AGTTCAAAAG	CCAGAAATAA	TEGCCETCTC
181621	GGATGGCTAA	AGTTGAGTAA	TAAGAAACTT	AAAAGGACTC	CTTTTTTTTT	TGCTTTAGAG
181681	TGCTATGGTT	TATGGTTAAA	AGCTTAATTA	AAAGTGGATA	TTCAATCTCT	AAAAGCCTGG
181741	GACTCCTTGG	GAAAAGCAGA	GGAGGCACCA	CAGACCCCAT	TTTGGGAAAA	CCTCTCTTTT
181801	CCTCATGAAA	CCCCAGGAAC	TGGAAGTGGA	TAGATCCTTC	GCAAAATCTA	AGGCTCTCTT
181861	TGGCTTTGCA	TTATGTTATC	TGATGTTTTT	GACTTTTGGG	GGTATCAGAA	ATTACTTCC
181921	ATTATGAGGG	AGATCTGGTG	TGTAATAACC	AGGTAGGAAA	TATACTTCTG	GGGATAGCTA
181981	AAGGCAAATA	TAGGTGAATA	CTTGGCTATT	TGCACTTTTG	GATCACAAGA	AGCATTCTCT
182041	TGACTACCTA	GAAGGTATGG	AAATGTCTCC	ATCCCCACCG	AGAGATAAGA	TTCCCAGGG
182101	AGATGGCTGA	TCCCCCAAAA	GAGGGCTGAT	TCCCTCTTTT	GGGATCCAGG	ATCTGGTATA
182161	AAAATGGGAC	CCTGGCCAGG	CACAGTGGCT	CACGCCTGTA	ATCTCAACAC	TTTGGGAAGC
182221	CTCAGAGTTA	TGAATGTCTC	ACCATACTGA	CACTTTGTGA	CTGAGCTCCT	CTCTACCCTC
182281	GACACAAGAG	ACCCTAATAA	TTAGACAGGA	ATATCATTGC	CCCTATTTAG	TCTGAAGAAG
182341	TTATAGAAGA	CGGATCTTTA	TCCCACTGCA	ATCCTTAGGA	TTAAGGGTTC	CCTCGTAAAA
182401	GGGAGTGGGA	AAATATGTCA	GAGGCATTTG	AATCAGAGTG	ACTCCATCTT	CANTACCCCC
182461	TGGGTAAAAT	AAGGCTGAGG	CCTGCTGGGT	TAGGTTAGGC	ATTCTAACCA	CCA CTTTA CT
182521	CACAGGATGA	GATAGAAGGT	TGCACAAGGT	ACCCGTCACA	AAGACCTTCC	TCATAAAA
182581	GGTAACGGTA	AAGAAGCCAG	CTAAAGCCCA	CCAAAACCAA	CATGGCCACA	AAACTCACCT
182641	CTTGTCATCC	TCACTGCTCA	TATACACTAA	TTATACTGCA	TTAGCATGCT	ACAAGIGACCI
182701	CCCACCAGTG	CCACGACAGT	TTACAAATAC	CATGACAACA	TCTGGACGTT	ACAAGACACI
182761	GTCTAAAACG	GGGAAGAACC	CTTAGTTCTG	GGAATTGTCC	ACCTCTTTCC	TCAAAAATTC
182821	TTGAATAATC	CATTAGTTTA	GCACATAATC	CAGAAATAAC	TATACGTCTC	CTTATTTCAC
182881	CAGTCCATAC	TGCTGCTCTG	CCTATGGAGT	AGCCATTCTT	TATACGICIG	TTTATTTGAG
182941	AGATAAAGAC	TCGCTCTGTC	ACTCAGGCTG	GAGTCTGGAG	TOCHTONO	TCTTTTCCCT
183001	CACTGCAACC	TTCACCTCCC	GGGTTCAAGC	AATTCTCCTC	CCTCACCCTC	CCARCTAGCT
183061	GGGACCACAG	GTGGGTGCCA	CCATGCCTGG	CTAATTCTCCTG	TATTATTATT	ACACCARCTAGCT
183121	TTTCGCCATG	TTGGCCAGGC	TEGTETCEAA	CTCCTCCCCT	CARCCCATCC	AGAGATGGGG
183181	GCCTCCCAAA	GTGCTAGGAT	TACAGGCATT	ACCCACTATG	CAAGCGAICC	ACTIGCCTTG
183241	CTTAACTTTT	TTTTGTTTTT	TTGAGACAGA	GTCTCACTAIG	CAIGACCCAI	CTACACCOMO
183301	GAGTGCAGTG	GTGCGATCTT	GGTTCACTGC	DACCTCTCCC	TCCTCCCTTCC	CIAGAGGCTG
183361	TCTGCCTCAG	TCTCCTGAGG	AGCTGGGACT	ACCICICC	CCCACTACAC	AAGCGATTCT
183421	TTGTATTTTT	AGTAGAGACA	GTGTGTTCCC	ACAGACAIGI	CCCTTCTCTC	CCAGCTAATT
183481	CCTCAAGTGG	TCTGCCTGCC	TCAGCCTCCC	ANDITIGICA	CAMMACACC	GAACTCCTAA
183541	GCGCTCGGCC	CTTCTTTACT	TENGLETCE	AAAGIGCIGI	ACTIACAGGC	ATAAATCACT
183601	CCCAAATTCC	TTCTTGTGTG	AGATCCAATA	ACTIGITITE	ACTITACTGT	ATGGACTAGC
183661	GCTGTTCAGG	CTGGAGCAAG	CTGGAGGTGA	TOCTOCTOCT	CIGIGAAAGA	ATGTATTGCT
183721	CTGTGATCCC	AGTAAGAGGA	TCATGGTCAC	TCCACCCTCA	ACCACACGAG	CATGCGTGAT
183781	CTGTAAGAAA	AAAAAATTAC	TAGAGGGGGTT	TARCACCIGA	TTTCACAGCAT	GATATCTCAT
183841	AATCAGTGAA	CTCAAAGATA	GGTCNATTCN	AACAGCAAA	TOTALCAGE	AAAAAGAAGT
183901	CAGAATGAAG	AAAAAGAAAT	DCACCCTTAC	AATGATCTAC	TOTGAAAAAC	AGAAAGAAGA
183961	TATGCATAAT	CCCACTCCTA	CARCACARA	AGACAGGGGA	TACCATCAAG	CATACTAATA
184021	AGAAATAATT	GGGACTCCTA	TOCOLOGRAM	AGTGAGAGGA	CAGGGAGAGA	GAATGTTTGG
184081	TTAGGAGGTC	TCTCAAAGCT	ANGENEGATA	GGCAAAAAAG	CATTAACTTG	CATACATATT
184141	ATAATCACAT	AATGAATTCC	AAGTAGGATA	CACTCAAAGA	GATCCATACC	TAGACACATC
184201	ATAATCAGAT	CNNAMACONNA	TADAAGAAGAT	GAATUTTGAG	AGCAGAAAGA	AAGGAACAAT
184261	TCATCACATA	CAAAIAGIAC	TOWNS	ICTGGAGTAG	GTATACTAAT	ATCAGACAAA
184321	ATAAACTTTA	AGRIAAGCAT	IGTTATAATA	AATAAAGAAA	GGTATTTTGT	AATGATAAAA
184381	GTGTCAATTC	ATCAAGAAAA	CATAACATTA	TAAACATACA	TGCACCTAAC	AACAGAGCCC
184441	TAATATTCAT	GAAACAAAAC	IGACAGAATT	GAAGGGAGAA	ATAGAAAATT	CGACAATAAT
184501	AGTTGGAGAC					
	CACTTGAAAA					
184561	GCAGAATAAA	CATCUTTUTG	AAGCTCACAT	GAAACATTTT	TCAGGATAGA	CTGTATATTA
184621	CTTCATGAAA	TAAGTCTCAA	TAAATGTAAA	AGGACTATAA	TAATAGAGTA	TATATTCTCT

Figure 8 (Page 57 of 73)

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194401	CAGGCATGAG	TAGTACGTCT	TGGAAGGTGT	GGTCTAAAGC	CTAGACTCCT	ATCTGCTTCC
194461	TTCAGCATTC	TCCAGTGTAT	CTGTCATCTG	TCTACCTTAG	GATAGGGGTC	TCCAGAACTT
194521	CCATTCACAT	TTAGAAGAGG	GCAGCGGCTT	TCTATGGAAA	ATATGAACTC	TCATTCATCT
194581	CTATTCCTTC	TTCTAGCTAT	GGTCCAGCTC	AGCTGTTTGG	AATAAAGTAT	CTATATGAAG
194641	TCTGCGAATG	GTTCTCAGAC	TGGTTGAACA	TTAGAATCAC	CTGAGTACCT	TCTAAAATTC
194701	TTATTACCCA	GGGCATATCT	CAGAATGAGT	ACCGCAGGGT	AGGGATAGGA	TTAGGGATCA
194761	TGATCTCTGG	AGTCTGGTTT	AGGCACTAGT	GCTGTTTAAA	ACTACGTTCA	TGAGGTGGAG
194821	GTTGCAGTGA	GCCGAGATGG	CGCCACTGCA	CTCCAACCTG	GGCGACAGAG	TGAGAGTCTG
194881	TCTCAACAAA	ACAAAACAAA	AAAAACCAAC	TACCCTTGTG	ATTTGAATGT	CCATCCAAAA
194941	TTGAGAACCA	TTAGGTAAGG	CCAAGCTGTA	TAATTAAAGA	GCAGTTTTCA	TTTGTCTGGT
195001	GTGGTGGCAG	CTTTTTGATA	AGGGAAGTAT	TGTTGCCATC	CACATACCTG	AGCCTCACTC
195061	CTGAGAACAC	TGGTGTGTAT	GTTGCTAAAA	TTCCCCAGGT	GATTCTGAGG	TTCCTTCCTG
195121	GATAAAAACC	ACTGACCCTG	GGAATGTACC	CACTGCCAAT	CTCCTGCGTA	AACCTTGGAT
195181	ACTGGGAAGC	CTACAGTTGA	AAATATTGGG	CTTGAGATCC	TGAAACAAAT	CTTGTATTTC
195241	ATTAAGACTA	ATATTTGGTA	CAGTGCAGCA	AATCAAGGGA	ATTTTGGTGG	CTGAGTTCTT
195301	TTAGAACTTT	TGCATTGAAA	TAGGTTCAAG	CAGCAATAAG	TTAAAACTAC	AACCTCAGCT
195361	AAAGGATTAA	AAGACACGTG	AGCTGGGTAG	GATGAGGTCT	AAGGTTGGGT	GTGGCGGCTC
195421	ATACCTGTAA	TCCCAGCACT	TTGGGAGACT	GAGGTGGGTG	GATCACTTGA	GGTCAGGAGT
195481				CCCATCTCTA		
195541	GCTGGGCGAG	GTGCCAGGCA	CCTGTAATCC	CAGCTACTGG	GGAGGCTGAG	GGAGGACAAT
195601				GAGCTGAGAT		
195661	TGGGTGACAG	AGCAAGACTC	CATTTAAAAA	AAAAATAATA	ATAATAACAA	TAATAATAAT
195721				TGGGGCAGAT		
195781				TTGGAGAAAT		
195841				TCTTGAGGGG		
195901				GTTACATAAA		
195961				GGGACCAGAG		
196021	CAGATATAAC	TAGCAGACTA	AACGGTCTAA	AAATAAAAAT	CATGCCCCAC	TCTAACCAI
196081				GTTTTTCTAC		
196141				CTGGCATATA		
196201				ATGTATTCTT		
196261				TACTATTCAT		
196321				GGCTTAAAGC		
196381				CTTAACTGGG		
196441				GAATTCTCAT		
196501				GAAAAATTCA		
196561				AGGCTGTCTG		
196621				CAGCCTTGAC		
196681				CACAAGTGTG		
196741	AAACAAACAA	ACCARARARA	ACCCCCACAC	AACTTTGTAG	IGCCATCACA	CCTGCCTAAA
196801	CCTCCCCCTCA	ACGAAAAAAA	CTCCCTTACC	CTAAAAGTTC	AGACAAGCTG	GTCTGGAACT
196861	ACCATACCTC	GCATATCCCA	A CTCTTCACC	AGGACAAATA	CACAMCAMM	GGTATAAGCC
196921				GTCCTCTATT		
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				TCAACAGTAG		
197041				TTCTTGGTGT		
197101	I I C I FAACCT	CITCITGCCC	TCTGGGGCCT	AAGATGAGGG	CTGTTATCAG	ATGTGAGTCT
197161				TCCGTTCAGC		
197221				CCAGATCACA		
197281				TGAGTGTTTA		
197341				CTTTTGGTGA		
197401				TTTTCATGTC		
197461				AGAAATACAA		
197521				CCCTCTTATC		
197581	GAACAAAAGT	CTGGCTTCAT	TCTATGACCC	CCACGTTGAG	TTTCTTAGTA	GCACTTACTT

Figure 8 (Page 61 of 73)

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AADATAAAAT	ACAACAGCAA	TOATTTAATO	STSTTSASA	TADDDAADTT	TOTTOTOTTT	705687
TAATAAATAA	ATAAATƏTTA	ADATTTDDAD	TOTTOADDIT	TTADDDAAAD	TDACTODOTT	TZ9ZOŹ
TTADTTODAT	AADDDTTDDA	ADADDDDDA	DDTDTDDDAA	ATOOOOTITI	ADDTAAADAA	T95202
AAADAAAADT	DDAADATDAD	TOOTOADOTA	AAAAA DD a B	AADDDDAAA D	DADDTDDAAT	TOSZOZ
TTODDTODTO	ADATTOTADA	ADTTDATAAA	AADADAATAA	DADTDADAAD	GAGACAGGCA	505441
AATATTOOTO	DDAADADTAA	TOTOATOOOD	DDTADADDTT	TTAADTOOTT	TTADADDDDT	T88Z0Z
ADADTTTTAA	DAATTDAATA	ATTTDTADAT	TTDABBBAAA	ADADTATATT	DDATTATATA	505357
AATOTOOAAD	ADDDDTAATD	DTDTADDTAT	CCTCCAAAGA	TOTTOTAADT	DDDATDDDAA	T97707
AADTDDDTAT	STSSSSATSS	TOOTTOTOAA	DTTTDTTAA Ð	DAADDAAADA	AATAAATTT T	TOZZOZ
TAABADDAAT	AADTAAATAA	TATDAATADT	ADADDDADTA	TTDDDTTADT	TTTATDAATA	SOSTAT
TTADDDDTDD	DODITOADDDA	DTADDDADAT	TADDDTCCT	AAA DDDTDDD	ADTODETODT	TBOZOZ
DTTABDBTAT	TODDDTOOTT	ADSTTOTOST	DDDAATDDTT	DTATOCOTAT	GGAGACGGGA	20202
TOTALTITIE	DOTTTTTAAT	DDAADTADAA	DADDDTDTAD	DDATATDADD	DTDDATDADT	T96T0Z
つつてつつてもなつも	AADTDDDDAD	DTDAADTTDD	DADDTDADTD	DADTOTAADD	TOOTOACOTO	501901
455T755AT7	SETTATOTOE	TTOTODOACA	DTOTTTOTTT	TTTTTTTTT	TCTTTCTTTC	201841
	PITTOTTOT	TTOTOTOT	DIDITIOITI	DTTTCTCTTTC	DIDITITIOT	187102
	OTTTOTOTTO	TTTOOTOTTO	AAABAADDBA	ATTAAATDDD	CATACCTTTT	TZLTOZ
TATAAAADAA	TTAAADSTAT	TOODICTOOL	ADTOBADTAT	DOTOAADAOO	ATOTTTTTOA	T99T0Z
T00 40T0T04	つつてょうううううび	TTSTTSTSAT	TTATAAATTA	TATTATTOUT	よようようせつうらら	TOPTOZ
GABTACTTEC	TTADSSTSTT	DATTOTAADT	DTDATDTAAA	AASTTTSASS	TATTAAADAA	TPSTOZ
クレンドエスロインので	TTAAAATATD	DADDITAAAAD	DTTDDADADA	DTDDDDTABA	ADAADTAAAA	201481
ンサイトをもといる	TATABBTBTT	AABAADDDTD	TOADADADAD	CACACCATAG	GATCATAGCA	707477
CAGACATTE	ATABBABADT	TTTDADTADA	ADTITADDAT	TOOOTOTITO	ADTATTATAA	TOETOZ
	DARAAADDDD	TOATOAATAT	DDTADTDDD	TOTITIONAUT	VOLONICATIV	TOTTOT
4724242544	TATTTOTTT	CTTCCTTGGG	AAATAAƏTTƏ	DDAADADDUT.	WONT COLUMN	TPZTOZ
	TADDATDDA	DADDADTDDT	DIDIDATITA	AASSSTSSSS	DAATITADAM I	TBTTOZ
DED4 ADDDED	DADDITITA	DATTOTTTAT	TTAAAAAADA	ADAAATATTO	VOLUCER 1997	SOTTST
ATANTATAAA	ADAATETSET	DADADTTDTD	TDATTOTOTO	ATSTSSSAA'T	TWILLINGE	TOOTOZ
DAAATDADTT	ADTTADTTAA	DAADDTAADD	AADTDDTDDT	TOTIGATIGE	ADADADATOO	TOOTOZ
ASSAATTEE	DODADTDADD	DTTADTADTA	AAADTATATT	DITIDAATTO	ADTOAATTIA	T7600Z
PTTDADTTAT	TODATATOTA	ATTTADTTTA	ATTICTITAA	DATTDTAAAT	ATTIDIADIT	Z00887
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207361	TCTCGAACAC	CTGACCTCAA	GTGATCCACC	CACCTCAGTC	TCCCAAAGTG	CTGGGATTAC
207421	AGGTGTGAGC	CACTGCACCC	GGCCGATACA	TGTGTTTTTA	AAGTCACAGA	AATTCACAT
207481	GTCTTGAAGG	ATTTTAAGCA	ATTTAAAAAA	TAAAGTCATA	GAAGCTTCAA	TTTAGGAATC
207541	AATGGAAAAT	TGATGATATT	CTTAGGATAT	GGATTTTTCC	TAAAAGAAAC	AAATGTATCC
207601	ATCCCCAAAG	ATAATTTGAT	TAGTATACAA	TTAAATT	AAACATGTCC	ATATTTACAC
207661	CCATGAATTC	TCTTTGCCTG	TCACAATAGC	TGGATTTATT	CACAATTGTA	GTAATTAGAG
207721	CCTGTTCATT	ATAATTTTCT	AGGTGATATG	AAGACTTTGT	CAGTCCAAGC	AAGTCTCCAC
207781	ATTGTGTGTA	GCAAACATGA	GAATAAACAT	TTTAAACTTT	TAAATGTAAT	AAGIGICCAC
207841	GTTATGTAAT	GTCATCCTTC	ATGTTCGAAG	GCACATGGAA	CATTGTTCTG	GTGGTDGDGD
207901	GGGGAGAGAA	ACACCATCAG	AATGAAAGGA	AAGACCGCTC	TGGAACCTTC	CTCCTTACCT
207961	CTTGAGCTTA	GTTTAATTGT	CCTGTCTTAT	GGTCTGCTAC	AAGCAATACC	ACTOTTONOO
208021	TTCGCATGCT	TCTCTGTGGT	TTGATAAAGT	ACATGCAATT	TTTCATTTAA	TTCTTCCACC
208081	TGCACTAAGA	AAGGAGCCTT	ATCTTTATTG	AACAGATGAG	GAAATGAATG	ATTACACAAC
208141	TTAAATGACT	AGCTCTAGGT	CACACAGCTG	GAACTTACAG	CCAGATTTCC	TTTTTACAGAAT
208201	CCTGTAACCA	AAAGCATACC	AGTAGTGCCC	CATABABATCT	A A CTTATA CA	COMOMOMO
208261	GTCAAAACTT	TTACTGATGC	TAAGAGGAGG	CARCATTRANC	AAGITATAGA	GCTGTGTTGG
208321	TTATGTTTTG	GATTATGTTC	TCTCCATAGA	TARARCACTO	MAGGGGAAAT	TATTTGTGTA
208381	GGCACAGGGA	AACTCCACCA	CAAAGCGTGG	TACCATTTCC	CLGIAGTAAA	AGAGATTCAG
208441	GAAGCCTGCC	ACCAGGAAAG	GTAAACCCAC	TECTOTO	CACAGAAGCT	AAATGGACGG
208501	TGAAGCTTAT	TCCGACACAT	TTACACACCTC	CTCCATCACA	TGCAGGCTAT	GTTAATAAGC
208561	TCCCAGTGTA	ACATTGGAGC	CACCACATCT	CCCCCATCACA	CTGACCCTTC	GTAAAGATAC
208621	ATGAAATCAT	CTGTGAGAAA	TTARCCCARA	TARGARECT	GTTGCTTTTT	CCTTAGCCCC
208681	GAATAAGTTT	TEGENANCEC	TTANGCCAAA	TAAGCAATAA	ATCCTGGGAT	CTAGGGAGTG
208741	CAGGCTGGAG	TGGGAAAGTC	CCAMCTOCCO	TITTTTTTGA	CTGAGTCTTG	CTCTGTCTCA
208801	TCATTCTCCT	TGCAGTGGTG	CGATCTCGGC	TCACTGCAAC	CTCTGCCTCC	CGGGTTCAAG
208861	CATCAATTCTCT	GCCTCAGCCT	CCCGAGTAGC	TIGGACTACA	GGCACACACC	ACCATGCCCA
208921	ATCTCCTCAC	TGTATTTTTA	GTAGAGATGG	AGTTTCGCCG	TGTTAGCCAG	GATGGTCTCG
208981	GGCCACCACC	CTCGTGATCC	ACCGGCCTCG	GCCTCCCAAA	GTGCTGGGAT	TACAGGCATG
209041	TAATATTCTC	CCTGGCCCGG	GAAAGTCATT	Traaaccaac	CTATGTATGA	ATCCCTACTA
209101	TAMIMITUIC IC	ACCAAGCGGC	TGGCTCTTTC	TCCTGAGCTT	GGAAACCTCC	AGTAAAATGG
209161	TTTCTTCCAT	TTCCCAGACC	ACCACTCTTA	TCTGTGAGCT	TTTTTGGCCA	ATTAAAAATTA
209221	CTTTTCTTCCAT	TATATTTTTA	TCTGTGTCTT	CACAGGTTTT	CTCTTTCTTT	CACTTTAGTG
209281	TTCCTCTCTC	AATAAGCAGG	AAAAATCCAA	TCTATCATGC	ACATGGGAAC	CCTTTCAATA
209341	TATATTAT	GTTGTTCCAT	TTTATGGGGA	TGCTTTTAAA	GAAAAAATTT	GTCCTTTCAA
209341	TATATTGAAT	ATCTTCCAGC	ACCACATCAC	CTGCAAGCTT	TGTAAAAATA	GTTCTACATA
	CLCARTTTTT	TTTTTTTTT	GAGATTGAGT	CTCATTCTGT	CACCCAGGCT	GGAGTACAGT
209461	GACATGATCT	TGGCTCATTG	CAACCTCTGC	CTCCTGGGTT	CAAGTGATTC	TCCTGACTCA
209521	GCCTCCCGAG	TAGCTGGGAT	TACAGGCATG	CATCACCATG	CCTGGGTAAT	TTTTGTATTT
209581 209641	TTAGTAGAGA	TGGGGTTTCA	CCATGTTGAC	CAGGCTGGTC	TCAAACTCCT	GACCTCAAGT
209701	COTTACTO	CCTTAGCCTC	CCAAAATGCT	GGGACTACAG	GCGTGAGCCA	CTGCACCCCA
209761	COLAGITITI	TTTTTTTTT	AAGTTGAACA	TATGTGAAGG	CAGGACCTAG	TGACACATAG
	CAATAACATT	TCCAAGTAGA	CATTACACTA	GGGAATTAGT	CGAAGTGCTC	ATTTAAAGTA
209821 209881	CCATCTCTCA	AATGTATTAA	AAGAGAATCC	TTGGATGTGC	AATACCTTAA	TTCAAAGGCA
	GCTCGTTATG	TATAAACTCT	CAAGCTTTGT	GATAAACAAA	TGTGCATAAC	AGATGGGACT
209941	ATTCACTTAC	AGCCCAGGGA	ATTTTATTGA	CGCTGAGAAG	GTTATGTGAC	TGGCTCTGCC
210001	ACTGTCATCC	CCATTCACTT	CATTTTGGAG	CAATATGACA	TAAATGCCTT	ACATGTGGGT
210061	TTTCTCTATT	TATCATGTGT	TTCCTATCCC	CTTGAAAGAT	GGCCATATTT	GCTTTACTTG
210121	GTTATAAGAT	CCCATATTCG	CTGTCTTGAA	GCCAACCAAA	TAATTTGACA	AAGTGGGTTT
210181	GTAGTGCTGG	CTATTTTGGT	GAAAAAAAGA	CAATGAGACT	TCATGTGTCA	TCCAAAGTTC
210241	TATCAGATCG	AGCTGTGAGA	GAAAGGAAAA	GAAAGGGGTC	TCAGTCAGGA	TGCTCACTAC
210301	ATACATCTGT	GTTGTTGTCT	AGGTCCAGAT	TTCTGTTCAT	TACGCTATGG	GCTGGCTCTT
210361	ATCATGCACT	TCTCAAACTT	CACCATGATA	ACGCAGCGTG	TGAGTCTGAG	CATTGCGATC
210421		TGAACACCAC				
210481		CCTTCAATAA				
210541	ATGATGGAAA	ATAGGGCTCT	TTGTTGAGAG	AAAAAACTTT	GAAAGGAAGG	CATAGATCTT

Figure 8 (Page 65 of 73)

Eigure 8 (Page 67 of 73)

DADADITIDA:	DOTADODAD	A SSSAASATS	r <mark>btəattaət</mark> t	AATTADDAA T	AADATDDTAT	12071
エンシエのみなすずつ	r ADAABBETA	A DAAADAAAD	r TADDAADATD) TTTDDADAAAT	TTDDAAAAT	T969TZ
OTATAATED	a papititato:	r TTOODTOOTA	AADAƏTTADA	DIDIDDDAG	ADDITATTO	106912
エンハナンのフェン	D STAADSTAT	D DADTAATTAA	A TOAAATTTDA	, DADDTTADTA	ATOTOAOAAT	T789TZ
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エタエンタシエンング	AADSTDASTA	A TOTATOTOO1	r TTDTTDDAAD	DIADITIADI	ADADDETDDD	TZL9TZ
THIPTITAAASE	TTADETEAM	r errorore	ATAATAAADA	TAADTTODDI	ATDTADDTAA	199917
エムつつてよるのだ	ATTTCTTTT (r AAAAADATƏ	r TATDDADDAA	TTDTADATDT	TTDDTTTTAT	109917
エムムエンシエムシ1	r AATTADTOT	r TODOTAADT	DESTORMENT	ATAAATTADT	DTDDTAAADD	T \$\$9TZ
DATATITITE	TTTTADITIO	A DATDITITAA	A DABAATAATA	ADDITAADDA	ATAAASTTST	T89T Z
TOTAGOATA	AATTTTTDTA	4 DADTDDTDA1	DAAAADAAAT	TTAĐĐĐĐAĐA	TTAATDADOT	77977
AASTSASTAA	DAAAATAAA	TTTDBATTT	r raproceres	TADTADTE	DDTTDADTAD	198917
エエエエンエ 4 エン ゴ	TADDIAADDA	A ADDDADTTDE) TAAAƏTƏTƏA	DATOCACTAG	DTTTTOOTOT	108912
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エゟつエエエ & デ & ^	: TTOTOTOTAL	X DTTTTADATE) AADTDATADA	AASTUTABAA	ATMICCAGILI.	190917
TOSTODSTOA	ADTOTADADO	ADDATOTOTA	ADADDIDITO	DADBAADTAT	ADDITION	100917
TOTOADOTAD) TETTOABOOA	(DDDADBDTAD	TGACTATTCA	TOADDADATO	りかみつかつもつかっ	176512
ASSEAGEST	. DITEBIEDIT) DTTDDADAAT	TADTODAADA	ATCTAGTCAC	WITITIWATE	188517
TODTTODOAA	DIDIAAAA A	TADTADAADD	TAAADTATTT	SOTOTAAAA	AAAAIAIAI	TZBSTZ
つかすつかけるつるこ	DOADDADATE) DADBETTTTA	CTGAGCATC	STTTASSTSS	SOMMITCIAMAGE	194512
ムエインテンエザイン	ASTBABTBBI	TTDTADATDT	CCTTTTATC	SSTORYSTICS	SERVINGE	TOLSTZ
441745555	DIBTBIDID) TADTADTTDD	TABBTTTADB	VD.I.SSSW.It.	2211111222	179512
すらずずずずずむ	AATADTTTTĀ	TADTTTDDDA	ACGAAGTCTC	DDDDVDDDV.I.	J.J.S.J. I WOWSO	TBSSTZ
られてられつるつてつ	DADDSTDTDA	, ATDTTAATAA	CTCATTGCCC	TOATAAATAA	HOWITING	TZSSTZ
SASTSTOTA A	AABBBBTDAT	TTDATTATAD	TATATTAATT	TTAATTUTT	WITHITHIE	194517
TOTOADADAD	TAADAADTTA	SOBABBABTT	CACATTGGTG	LYCHILLCCL	TTWOODWRTS	109512
AAADADTDTD	ATTDABABAD	DATADTDTDD	DOTAATOOTA	AAAbaaabut.	WHISTOWNS	176512
TAASAADSTA	DIABATTATA	、コエココロAエコココ	OATTTECCTGC	A LAMOTOTITIE	TTTTTVOODY	TBZSTZ
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TTABOTABOA	DIDATOADIT	TACATGGCAT	DDDAAADDTTT.	CHARTETACAC	TTTTTOTTOT	TSTSTZ
TOTOTOTATA	4 4 D T D T D T D T	DDATATDATT	AATGCATTTC	TYCCCTOLYT	INTRICTOR	TOTSTZ
つかつかららずずるて	ADSTATTTET	DAADATDTAT	TGTGTGTTG	D.T.W.T.W.D.D.D.W.D.	TTTOTIVITY	TROSTZ
4 45 7 45 7 7 4 5	DTATAAATTD	AAATƏƏTAƏA	DOTTADDTDA	PARTICITED	OH I WHOW I WU	186917
SATABLE	AAATTTTATT	ADTAAAATTD	CATTGAGCGG	DIAMMATA T	DAU LDAL LAS	126712
22455577775	ADTITIAAADT	ATCTTTTTT	DATTTTATAT	LIWITHITHIT	TINVINITUS	79877
422242247	ATATABADTD	DTATTTAADT	ATTATTAATA	LIDIAMATA	TTOWANDO	108412
22223333	೧೧೯೯೯೮೦೦೦	ADATTAĐĐĐT	DDTDAAA DDD	TO SECULA	2226221461	274741
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	<i>AAAAA</i> TOTAT	エつうATAつうエゴ	ATODIATOR	2211WWW222	0101100	STUUET
	コート スペインシーン	つつエダエダコニアシ	ひょうりつつ まつの	121200121	22112122	786777
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2721100010	これがらずがつり 4.4	222755TTA	つつりかんしつしょう	WWWATER	TW00	214261
	AAATDADADA	ナインのエムンエエン	D.I.I.D.I.D.I.WWn	TTTTWWDWDT	* > 0 > > > + + + + + + + + + + + + + + +	70277
	つつてんりるるるでる	AAATOOTTOO	ATTOAAUUAA	WWY TWY TO T T	271224271	214141
		つエつエムシエンシエ	OTATOATAAA	DOLLINGOLD	21274224	T8077Z
		エインエエシンシエン	DIDDOMEST	Wataawtttt	2404201001	274027
-000004040	A T つ A つ つ T シ シ A	TOOTOADTUT.	J.つりかつつ t かつつ	TIMMUTAN	C112C11C1	196612
	りょうかつりてわつて	DIDDEADDUD	.T.D.T.J.D.T DO T T	WATERTAND	24121222	773907
OTT ATOOTAD	TOTATOOTAO	DTTTDDDTDA	DATODIATOA	DTAADTDTTA	CABASBABAS	273847

22222						
220321	AGGGCTTTCT	GAGGAGGGTC	ACACAGAAGA	CCAAAGAGAA	CTCATGTTGA	ATTGAGATGG
220381	GITGTAGTGA	TAGTTGTCAA	CAGCCAATAC	AGAAACAAAA	AAAAACAAAA	CAAACAGCAA
220441	CAACAACAAC	AAAAAAAAA	AGAGAAGACA	CAAACACAAT	GCCACAATGC	CATTTTAGGC
220501	ATAATTTTAA	ATGAGTAATA	TTATATGTTG	AAATCCAAAT	TTTCAGAAAA	ACATTAGTGT
220561	ATTTTATTT	TGTTTAAAGA	AATAACCATC	TCAACTCAGA	ACCCCATGTG	CATTTTGGCC
220621	ATTITGTTTC	CAATAGTTTC	ATAAACTTTC	TTAAGTAACT	ACTGCACATT	GTTCCTTATA
220681	TTCCTTGTGA	TCAACATTGC	AATACACAAC	TGGGAGGGCT	ACTAGAACTG	GTGTAGAAGG
220741	AACTTGTGAG	ATTGATCATT	TTCTCTGTTT	TTTACATCTA	GGATTTTGAG	TCTGGTTGGA
220801	GGAATGTCTT	TTTCCTGTCT	GCTGCAGTCA	ACATGTTTGG	CCTGGTCTTT	TACCTCACGT
220861	TTGGACAAGC	AGAACTTCAA	GACTGGGCCA	AAGAGAGGAC	CCTTACCCGC	CTCTGAGGAC
220921	ATAAAGTTAC	AAACTTAAAT	GTGGTACTGA	GCATGAACTT	TTTAAACATT	TTTTACTTCT
220981	CTCCATATTC	CTGACCATAG	ACTCAGCAGT	TCTTAACTCT	GGCTGTGTGT	TAGTCTTCCC
221041	TGGGGAGCCT	TTATAAGACA	CTGATACTTG	GGACCCACTC	CAGAGATTCT	GAATGAATTG
221101	GTCTGGGGTG	GAACCCAGAT	ACTACTAATT	TTTAGATACT	CCTTAGAGGT	TTCTAGCATG
221161	CGCCCGGGGT	TGACAACAGC	TGGACAAACT	TGAAAAGTCA	ATTCATGTGG	CCTTTGAATT
221221	TTCCTCATTG	GAAAGTACTA	AATAAATAAA	AATTCATGTG	AAAATGATCA	CTGATAAATA
221281	TCTTCATGGT	GGGGCAGGTT	ATTGGATGCA	GAGAAGATCT	GCTCGGAATT	GTAGCCATAT
221341	GTTACAGATC	TCAGCACCGA	TCGGAACTGT	AAAGCTATAA	TCCCCAGAAT	TAAAGTTTTT
221401	ATTATTTTTT	ATACATTGTA	AAACATAGAC	GTTTATTTAT	GTGATTAAAT	TCTATTAAAA
221461	TTTACATGCT	TAAAATAAAA	AGACCATTTT	CAAATTATTT	AGATCCAGAT	ATTTCCATCA
221521	GATTAAACAG	ATATTTATTT	ATCCTAGCCC	AATTGCAAGA	GATTAATGAT	GAGAAAATGA
221581	CCAATACAAG	AAATAAATTA	TGAGGTTAAC	TTAGAAATCA	AGGACAGAGA	AGATAGAACT
221641	GGAAGGCTTG	TATTGTGAGA	AGAATGAATG	TGAAGGAAGG	CAATGTAGAC	ACTTCCAGAA
221701					GAGAGATGAC	
221761					AAGACATAAC	
221821					TCACATACAA	
221881					TTTCTTACTT	
221941					CTCCTAACAC	
222001					TGCCACAACA	
222061					TTGGACAAGA	
222121					GGTATTTTTG	
222181					ATGAGAACAT	
222241					GAAAACTAAA	
222301					ATGTACAGAA	
222361					AAAATGGGCC	
222421					ACATATAAAA	
222481	CTTCATTAGT	CATTACAGAA	ATGAAAATCA	AAACTACAAT	GAAATACCAC	מדדממממדמ
222541					GCCAGACATG	
222601	GGAACTTTCA	TACGTTACGA	ATGTGAACTT	TGGAAAGCTG	CTCGGCAATA	TCTCCTAAAG
222661					AATGCACATA	
222721	AAACATGTAC	AACAATGTTC	ATAGGAGCAC	TATCTGTAAT	AGCCTGAACA	GGAAGTTGTC
222781	TGTTAAAAA	AGAATGAGTA	AATAAACCAC	GGTCTATTTG	TATAGCAATG	DGAATTAACA
222841					ATATTGATTA	
222901					AAACAGCTAC	
222961					TGGGGGTTGT	
223021					TATTTCTTGA	
223081					GTTTATGAAT	
223141					TCTGCTCTGA	
223201					CTTTGCTTTT	
223261					GGAGTGCAGT	
223321					TCCTGCCTCA	
223381					TTTTTGTATT	
223441					CTGACCTCGT	
223501					CCGCGCCCGG	
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Figure 8 (Page 69 of 73)

Figure 8 (Page 71 of 73)

					DAAADDTATT	186622
	TAAATAATAA					126622
TOOBABABTO	DTDADDOTTA	TDADADTATT	TASTTTTOOO	TOATTDAADA	DTTTDADTAA	198677
DADATAATDA	DTDDDADAD	DTDTDTTTAT	TTTOOTTATO	TTDTATATAT	DTADDDDAAA	T086ZZ
ATDDDTTTAD	ADTADAAAAD	TTADTDAĐĐA	AGCAGGCAAG	ATTAAAAADA	AAAATOOTAA	77627
ADDDTDDADD	TATATTƏTA	DTDAAATDAA	TAAATAADAD	AATAADADAA	ASTSTADODS	T8962Z
TOOODAAODT	AAADAĐADAA	ATDADDADAT	ADDDATODTD	ACCACTCTAG	Debadoetat	129622
DODDADDADA	DADTATDTAT	ATAAADDTDA	DOTODATTTT	DASTSSTDAA	TOODATADOT	195622
DAAAADADTD	TODOTOTTOT	DTDDATA DTD	ACAATGACAA	TODATATOOT	ADADTDDDTD	T056ZZ
TODOTOTITI	TTTOADDDAD	TTTDATATAT	AATTƏTTTƏT	TOTITITGT	TATTAĐĐĐTA	779447
DADTTDDDDT	DOTATATTAT	TTTDDTTDTA	TODAAAAAA	DOTTOOTOTA	CATGCTCTCC	188622
TTDDAATTOT	DATTTDTDAT	TTDAADTDTT	DDTTAATAAT	DTAAADATA	TTTTTAATDD	725622
TTTATOTTTA	TOAATTDAAA	DTTDTTTATA	TADTTATTAD	TADDATABAD	TCCTCTTTG	T92622
TTTDTDAATD	ADTTAAADTA	ADATODAATO	AADATTTATD	ATAADTOODD	TOOTAAACOT	T02622
TODATADADT	DTDTDAAATD	AADDADADAT	ADAASTATTD	TOTOOOSTIT	SSSSSSAAAT	771677
DIADDITDDA	DADATADDDT	DOTOBOTODA	TADDDTDTDA	GCCATTTTAG	ATTTDTDADT	180622
AAƏTAƏTƏTY	ADDDAADDTT	DTTDADDTTD	TTTTTAATOT	DEASTADDAD	DDTDTAADDD	120622
DADDTAAAA D	TADDADDADD	DADTAATDTT	TDAADATTTA	TTADTADDAD	ADDTDADTAA	196822
AADTADTADD	DDDDDADADA	ACAGTGAAGC	DEPARSEA	ATADAĐĐTAD	DDTDDTTAAA	106822
DADADTATDA	Abbbbbbtcc	DTATDDDAAT	DTADADDDAA	DTATTDDDDA	CAGTOSTO	T \$88ZZ
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233521	TGTTTTCTTC	CATGACTAAT	ACTTTTGTTA	TTATAGCTAA	AACTTCATTG	GGGCCAAATC
233581	TTAGATCATG	TAAATTTTCT	TCTATATTTT	ATTCTAAAAG	CTTGTAATGT	TTGATACATT
233641	CTAAAAGATG	TAATGTTTGA	TACATTACAT	CTAGTCCTTT	GATTTATTTT	TAGTTACTTT
233701	TGTATAAGGT	GTGAGAGATG	TCTCCAGTTT	CACTTTATTA	ACACATTGTG	GTGTTCCAGT
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Figure 8 (Pag 73 of 73)

Figure 9 (Page 2 of 74)

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10141		TTCATACCTC				TGTGTAGTTT
10201		TAACTAAGAG				CCTATAAGTA
10261		GCTCTTTACT				
10321		CCGCCCCGAA				
10381		AGCGCAAGCG				
10441		TCCATCCCGA				
10501		ACATATTTGA				
10561		CCATCACCTC				
10621		AGCACGCCGT				
10681		TCCAAGTAAG				
10741		ACTAAAAGAG				
10801	ATTAGAATGT	AGGAACTGGA	GAGGGGTGGG	GACAAGTGTT	GCAGCTTAGA	GAGGGACAAA
10861	GGGTCCTGAA	CCCGAAAGAA	GCCAGCCATT	AAAAATGGGT	TTGGGGTCAA	TTCGTTGTGC
10921	TTAAATTTAA	AATGGGGACA	AGCGGCCATT	TTGCTAACTC	GGCGTTCCCG	GAAGAAACCG
10981	CAGGCTCGCT	TAGGTTTCAG	ACCCAGCTGT	CTGTCCCTGT	CTACGTCGCC	AGGATCAACG
11041	GTTGCCGTAA	TGTCATAATT	TCGCCACCAG	CTTCTAGCCA	ATAGGCTGTC	CTGTCATTTT
11101		CAATCGAGGG				
11161	TGGGAACCTG	GGCAGTAACT	GCCTAAGGAA	GGACTCCCCC	TCTGTTTTCG	TGGCGCACAC
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11281		TGCGTCAGTT				
11341		AGATGTTGCT				
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11461	TGATTGAAAC	TTAAAATCTC	CGTAGGGGGC	TTGTAACATG	CAGAAAAGTT	TGAAAGTTGC
11521	TTTAGGAGAA	GCCAACTCTT	AACTGCTGGG	TAAATTGACA	AGCCTTCGAA	CACTGAACTG
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11641	AGCACATACA	CTGTGTCTCC	TAGAGGACTC	TCCCTTCCTA	GACAACTGCA	GGCCGCTTTG
11701		AATTCCACAT				
11761	TTTTAAGATG	AAGGGTTAGA	CGTAGTCTAC	CTATCTTTTT	ATTCAAGTCT	AGAACACGTT
11821	TTTAGCACCT	AGAAGTTTGC	TTTCTCCATT	AAAAACCGGG	AATATACAAT	TTAAAATTAAA
11881	AGTGTTAAAG	CAGATTTTTA	CAAACTTAAA	TACCATGTAA	TTTAGGTTAC	AGTTACTTAA
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Figure 9 (Page 4 of 74)

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22681	AGATGCCTTT	ATTTTATTCA	CTCACACACA	TATGTAGAAA	GAGAAATATA	TGGTAAACAT
22741	TAAAAAAAAC	AAATTAGAAT	GTAAAATTAA	TACTTTAAAA	AATGGGCTGT	ATACTTTTCT
22801	TATCACCGGA	GATAAGAATT	TATTATTTTT	AAAATAAAGT	TATTTTCTCT	GTGACTGTTT
22861	CCATGACTTT	GCTACTTAGA	AGTTAGAGAT	GCCAAAGTTT	ATCTAAGAAA	ATGTTTATGG
22921	AAATATTATT	TCAATAATGA	ATGTTTAGAA	GACTGAATTT	CCTGACTGGG	CACAGTGGCT
22981	CATGCCTGTA	ATCCCAGCAC	TTTGAGAGGC	TGAAGAAGGA	GGATCGCTTG	AGTCCGGGAG
23041	TTCAAGAGCA	TCCTGGGCAA	CACAGCGAGA	CCCTGCAGCA	AAGTAAAAAG	AAAAAAGAAT
23101			TCCTTTGGGC			
23161			CCTACATTTT			
23221			TATATACTAT			
23281	ACCATTTTAG	CTATCTAATG	CAAAATATGA	ATCTTTTTTT	TCTGGGTCTT	AAATTATGGA
23341			CTCTAATTCT			
23401			AGGATCAACC			
23461			GAAATTATCA			
23521			TGCACCTGTA			
23581			GAGTTGTGGA			
23641			ACAGTAGAAT			
23701			AAAATTTTTT			
23761			ATCATGCTCA			
23821			GAGTAACTGG			
23881			AGGGTCTTGC			
23941			CCTCAGCCTC			
24001			GAATTTTTT			
24061	TGTATGGGTA	TAACAGAGAG	ACAGAGAGAA	AGAAACTTTT	CTATCACACT	TTGCAATCAG
24121			TGGCTTTTGT			
24181	CTTTACCACA	CTGTCCCCTT	AGGCAAGGTC	TTTGCCATTC	TTCTGAGACT	ATTGCAACAG
24241	ACTCCCAACT	TCTGACTGTG	GGCCCTTCTC	AAAAATGATT	GTTTATGCAA	TAAATCTAAA
24301			ACAACAAATT			
24361			ATTCCCAGCA			
24421			GCCTGGCCAA			
24481			GTGGTGGGCG			
24541			TGGGAGGTGG			
24601	CACTCCAGCC	TGGGCAACAA	GAGCAAAACT	CTGTCTCAAA	CCAAACCAAA	ACAAAACTTC
24661			ACACAAGTAT			
24721			TATGCTCTGG			
24781	GGCCTCAGCC	TTTGTGAGCA	AGCTCTTATC	TCCAGGCCTC	TCACAAAGAC	CTGTTCCAGT
24841	AGAAGCTCAG	GGGAGCACAC	TGGACATTAT	TCCAACAACC	CTTTCCCCAC	AGCTATGCAG
24901			AATTAATTAA			
24961			TCAAGCTCCT			
25021			TGTGCCACCA			
25081	GGACCAGGCC	AACCTAGTCT	TGAACTCCTG	GCCTCCAGCC	TTCCGAAGTG	CTGTAATTAC
25141	AGGCATGAAT	CACTGCGCCC	AGCCAACCCG	CCCAGTCTTG	TTAGACATGG	GGTCTGTAGT
25201	TTCTAGTAGG	TTCTTGAGTC	TAGGGTTCCT	ACCTCATGTT	TTATAGTTAA	TTTAGGGGAG
25261	GGACTGTGTC	TGTTTATCTG	GGGATGTAGG	GGTGGGCAGG	GGGATAGAGG	GGACTTCAAT
25321	TAATGAAACC	AGAAGCAAAA	CTCAGTTGAG	GACACCGGTC	ATGAGAGTGG	CCTGATTATG
25381	GCCAATCTTA	CATAATGTGT	GAGATCTTGA	TATTACCCCA	TCCTTGAGAG	TCCTCTATAA
25441	AGCTACAGGG	ACTTGGGAGC	ACCTTTAATT	ACAGACAACC	CATGTTCCTG	TGGATTATGA
25501	TTTATTATT	TGCACATGCC	TAAATAAAGA	CATCCTCTGC	AGTCTTTTGA	CAATTCTATA
25561	DGCDTCTTCT	GACTCCGCAA	TTAGACAGCT	AAGAGATCTG	TGTTACTTCC	CTCACATATA
25621	דיייית מיי ממיי	AAAATAAAT	TCATGGCGTG	AATAATTTCT	TTCCTCTACC	GATTTGAAGC
25681	TATCCATTTC	GAAGACCACT	CTGAAGAGAT	GAAATAAGTO	TTCTGCCAAA	GATTACTTAT
25741	TAATTTACAA	GGAAAAGGGG	AAGTTTTGTT	CCTCTCCGTG	AATTTGATTG	AAAATCGAGG
25801	GCTTTCTCGA	ATAGTTTTG	CATCCAGGGT	CATTTTTCAT	' TAAAAAGAGA	AAAGTCATGT
25861	CAAATATGAA	TTTCCGCAGA	TTATTCAGCA	CTAGACCCTG	GGAGATTCTG	TAAAGAGGGG

Figure 9 (Page 8 of 74)

Figure 9 (Page 10 of 74)

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35701	AATAATTCCC	TARRARATIA	AATTAAGATT	CAGAACACAG	CCTAATATCT	AGTAAGTAAT
35761	TTCCTACATE	A A TA TA TA TOTAL	TTCTTAAGAT	TATATATATT	CATGGGGTAC	AAGTACAATT
35821	AAAACTT	AAIAIAIIGC	ATTGTGGTGA	AATCAGGGCC	TTCAATCCAT	CCCGGAAAAA
35881	ACAAGIIIII	GAAAAGATTT	CTGCCATGGA	AAACTTTTAA	TGTACAAATT	CATCCATCCA
35941	TARACARONA	AATATATAAG	TATCAACTCC	AAATCCACCA	TATCTATCTC	TTCTGCACCT
36001	AAACAATTA	CTCAGAAATA	GAATGCTTGA	GATACCAGAA	TGCATGCATA	TCAAGTAATA
	MAIGCAIGCA	GGATGTCAAC	GCATCCTAGG	CTTTCAAATA	AAATTGTCAT	ACAAAATACT
36061	TTAATATTGT	AGTAACATTC	TACATGTTAG	AGTGTAGAAG	TTAATCGCTG	ATGCAAAAA
36121	GGAAAAGAAC	ACATTATACC	CAAAGCCTAC	AGAGAGAATC	ACAATTACAA	ATATCAGCCT
36181	GCATGTGAAA	ATCTTTAATT	TGAAAGTCAG	AAATATTTAA	ATGATAGTCA	TTGTTAAATC
36241	AGATTGTGGT	TTGAAAAAA	GTTAGTTTAA	AACTGAGTTT	ATGAAAAATT	TGGGGATTTT
36301	AGAGACAGTG	TTTTGTTTTT	AAATGTGTGT	GAGTTTGTGA	AGAATGTTTT	ATAAAATACT
36361	GACAGTATTA	TAAGATGACA	TTATTATAAT	ACAACATAAG	AATTTTGGCC	TGTACCTCTC
36421	AGCAGTCCTC	AATCACCTGC	TGTACTTGAC	TCAATGATTA	TCAGAGTGGT	TTGTTTTCCT
36481	TCTGTTGTGT	TCCCAGTTCA	GGCAGCTCAG	CAATGGCCTG	TGATTCCAGC	AATTCAAATA
36541	GCTGGTAAGT	AGTTTCTTGT	TTGTTTTCTC	AAATTTTCAG	GGGCTTTTCT	CTACAAGTGA
36601	TTTCCAGTGC	ACGCCCCTCC	ACCCATTCTT	TATTCCTTTA	CCTTCAGGAA	AACCCTCAGC
36661	GCTGCATCTC	TGGTCACCGG	ACCACCGTGG	TACATTTACC	TATGGCCACC	AGGTGTCACC
36721	CTTCTCTTTA	CTACCATGGT	TTGTGAATGG	TTTTGCCAGA	GGTGAATAAG	AATTTAAAAT
36781	GCAGGTCTTT	GATTTTTCAA	ATGTAGTTGA	CCTTAAGAAT	TTATGAATAA	AGCCAGAAAA
36841	ATTAAGCTTA	AAAAACACCG	AAAGAAAATG	AGGACTTAAA	ATTTCTATTA	AAAAAATTAA
36901	CAGGCCACAG	TTGCTGATGT	TTAGTAAATG	TGTTAGTGAA	ATGTGTTACT	GTGAAGACTG
36961	GGGTGTTTCT	TGAAATCTCA	GCCCAGGTGA	AATAAAACCA	ATATAAAACA	AATGCTTACC
37021	TAATAAATTA	ATTGTAACAT	ATTCCTTATG	AGGTAGAAGA	GTAAGTGAAG	CCTTATAGCA
37081	GTCTGCTTTC	AGTATAGTAA	GATATTAAGA	GAGAAATAAT	TTGTCATATG	CTTTCAGAAT
37141				AACTTAGACG		
37201				ATGAAAATAT		
37261	GAAATTAGAA	GCAAAATAAA	TGTCTCCAAA	ATGACAAAGC	GATTAAGTAT	ATACACAAGA
37321				ATACAATACA		
37381				AGGTAATAGA		
37441	ACTGGCTAAT	TAAATAACAG	CTTTAATTGT	ATTCATTTTA	TAGCTTTTCT	ACAATGAGCG
37501	TAAATCACAT	TTACTTTTTT	CTACATAACT	TTTCTAACCA	CAAAAAAAGA	AAATGGTTTA
37561	AAAGAAGAGA	TGAGATATCT	TTGCTAAAAT	TTAATGCCTA	AAGAAGAAAC	TTCTGAGCTG
37621				AGACAGAATG		
37681				CATGCCATCT		
37741	TTTTGACGGC	TGGGCAGGGT	GGCTCACACC	TGTAATCTCA	GCACTTTGGG	AGGCCGAGGC
37801				ACCAGCCTGG		
37861				ATGGTGGTGC		
37921				ACCTGGGAGG		
37981				TAGAGTCTCA		
38041	TTGAACATGG	TGAACTGATT	TCCCAGAATC	TAGCAATTCC	TGAATGTCCT	GGTTAGATTT
38101				CTCCATGGAA		
38161				AAATGAGAGA		
38221	GGAAATACCA	CCAGAGTTCT	GACTCCAGAG	GCACTGGCCT	AGGENGENCE	CCCTCTCTCA
38281				AATTACAGTC		
38341				AGGGAGTCAC		
38401				ATCAGTTGGT		
38461				CAATGAGACA		
38521				CTGAATGTAG		
38581				TTCTATCACT		
38641				TTATTTTTAT		
38701				GTGGCGCAAA		
38761				CAGCCTTCCG		
38821				TTTTTATTAG		
	LUCCACCA	COCCCNOCIA	ALLILLIGIA	TITIMITAG	ACATOCOTT	TCACCATGII

Figure 9 (Page 12 of 74)

Figure 9 (Page 14 of 74)

			DDAAADTATA			10850
			TOTTOTADIO			4254 1
TTOTTTTT	DADTOCOTT	DASTEASADO	DOBADTETEA	ADTOBTOTOT	DTACTTCATG	T815+
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TDAAADDDTD	TACASTSSAT	DATOTOTITE	ASSACCAGGA	つててつしんもつつつ	TTTTTTCAA	T8877
TACTTGTTTA	DADBADDDDT	DATOTOODET	TOTTOATOOA	TGAGCCCATT	CCTCCTTC	44851
STATTTOTT	DAADADETDD	DODDADATOD	CACACCACAG	TTAASATSS	TTDDTDATTD	19400
TOADTADADA	DTDTDDAAA	ATTTTTAAD	GCAGCCAGAG	ATACACATA	DOTOTTAOTT	10400
CONTROLL	DATTTTDDTD	SOTSESSIVE	CAGCTTCCTG	AATOTOTOAO	DICOUNT	T 7 9 7 7
DATDADDTDA	SSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSS	COTTOTOTIT	TOOOTOAOTO	TTTDDTAADA	DADDADTTAT	T85**
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DDAADTDDDD	TODITIOAADI	DOBADETDAD	TODDIAOTDA	DADDOTDADA	TOAADTDOOA	187**
TODADTDTDT	DETTOTEDEA	CAGGGGAGAC	TDDDDDAOTT	TCTTCTCTTT	CAACTTTAAC	44221
TOAAADAADA	AATAAAATTA	AAATAADDDD	ADTOBTOADD	CCAATTACOO	TTADSSTAAA	19177
			AAADDDDATT			TOTOT
DTADAATTDA	AADTDTAADD	TOADTOODAA	DOTTOADOAD	TOODOTOTAT	SSSTSAASTS	10000
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DADDBADTDD	DIDDATOTAD	TAAADTDDAD	TODTDAAADT	STATTOAAS	SESTIBLIASS	12664
ADTTTĐĐĐĐT	ADABATBATT	TTTATOTTT	TAATDDDADD	CGCCACCACA	DTDDDDATAT	T98E1
DABBBTDDAA	AADTOOTOO	AATOOATOOT	DTTABBBAAD	TTSSSTOOTS	SASSTSSAAS	T08E7
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Abaccccaga	DADTDDDDDA	CGTGGCTGCC	ADADDDTDTD	TTTTOSTSTS		T05E4
			SOTEASETTE			7955
TOUTOTOT	DDADTADDAT	ADTDAADTAD	DTDDADADAD	OTODDADTTD	TADDADDADD	18562
DDADDATDAT	DTDTACCATGTG	TOODDAOAAO	GGAGACAGAG	TOOAASTETE	ADADADOTOA	43357
TODADAADTA	DODADBADT	STOSTOSTSS	GAAAGACCAA	TTCAAACAAG	SEARASSSEA	43261
TODADOTADT	DDTTDDADAAD	CATAGAACAC	TADTADTDTD	CACACGGCGA	OTOBBBBOTT	43201
DADDTADDDA	AADDTDATDT	TAAAƏTTƏƏƏ	TOTADADDDA	SOTOBACTOA	STOTOABAD'T	19169
TTCCCACCCTT	ASTSTADAAS	AGTCAACAGT	ADADOTDATA	ATTTTCCATA	DIDILIDATAL	43087
DDTTDDDATA	DADDTDTDD	DODDADTDDA	ADDADTDDAA	DADTOTODAA	AADTADAAAA	43057
DTADDTADAA	DTTDDTTTDT	DDADDTTTDAD	CAAGACCTCA	ADDDTTDDTTDD	OTTOTOTOTI.	T96Z#
DOTTAAADDA	ADDADTOTAD	TAAADDADDA	ACCATCCCCA	CCTATTATOOO	TODATAADDA	42901
AADDAADDTD	AATTƏTTƏƏ	ADDDDTAAAA	ADDITCTGGA	STSTSSSASS	TOOASTASSA	42841
つつつむADADIT	TADSTITITI	TOTTOTOATO	DOTOODATO	CGTACTTCCA	SSATASTASS	18724
SOTSSTEETS	DAADTTAADA	DDDDADTDTD	TOADOTAADO	TOTOSTOSTO	TTDDDTDDD	72724
DADDDDDTDD	TTOBADOTDA	TGCCTTTCCC	CCGAATCTTG	TCTGCCTGGC	TOTOCOOR	T9977
DDADDTDTDD	STDASDDDAS	DTOTODACOT	CAGCAACTGC	DYDDADDDOT	DTDDADAAAA	45601
STEETTELL	DOATADDADD	TTTDTDTDAA	DADDDDDATA	TOADATDADD	DIDIDADDID	T \$ 5 Z \$
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DADDTTTAAD	ADDDTTDDDT	DAAADADTTA	DDATCTCAGG	DATDDAATAT	TAADTOTADA	T Z 7 Z 7
TACTCTCAT	ATGACTTGTG	DDDAAATTAA	GGAGAAAAGA	TADSTATADD	DADTTTUUU	45361
TOADADATAD	AAGCAAGGGG	AAATAAATTD	DTTATTADAA	AADATATAAA	TATDUADAG	45301
TTTTAADTDT	TTADADTTDD	DDDDDDDDD	ADDBABTBBB	DADATTABBD	TOPLOVV	
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48601	CGGAGGTTGC	AGTGAGCTAA	GATCGTGCCA	TCGCACTCCA	GCATGGGACA	CARCACOANO
48661	ACTTCATCTC	AAAAAAAAA	AATTAGCTGG	GTGTGGTGGC	ATCCACCTCT	AAGAGCAAG
48721	ACTCGGGAAG	CTGAGACAGG	AGAATCGCTT	GAACCTGGGA	GCCGGACCTT	CTCCTCAGCT
48781	GAGATCATGC	CATTGCACTC	CAGCCTGGGC	AACAAGAGCG	A A A CTCCCTC	TCARAGCC
48841	AATAAATAA	ATAAAATGCA	DTAATTAAAA	GATTTTAGTA	TATTTACACA	CARARATAA
48901	CATTACCAAA	ATTTTACATT	TCTATCTCCC	CAAAAAGAAA	CCATCTTCCC	GATGTGCAAC
48961	ACCCTTAATT	CATCGCCTCC	CAGATTCCTC	CATTCTCCTC	CTCCTCCCC	CTAATTCAGT
49021	GACAATCTTT	AATCTACTTT	CAGAIICCIC	GGAACATTTA	CICCICCCT	CCCAGCCCTA
49081	TATATTGCTT	TGCCGTGACT	CITICIATII	ATTTAGCATA	GIATACATAG	AGGCATATAA
49141	CATGGACCAA	TAATATCTAT	TATARGACA	TACCACAACA	TATTTTTAT	GTATGTTTT
49201	CAGCCGATGG	ACATTGGTTT	CTTTCTACTT	TATGGCTATT	TATTITATT	ATTCATTCAT
49261	CATTTATCTA	CANCIPPRINT	TCTACACTTA	TGTTTTGATT	GGGAATAGTG	CTGTTATAAA
49321	ARTITATOTA	CAAGIIIII	DECCERROR	TGTTTTGATT	TCTTTTGGTT	ATATATCTAG
49381	TTTTCCAAAC	TA A CC A TITTE	ATGGTAACAC	TGTTTAACCT	TTTGAGGAAT	TGCCACATTC
49441	CATCTTTCCC	TAAGCATTTT	ATCCTCCTAT	CAGCAGTGTA	TGAGAGTTCT	GATTTCTCTC
49501	TOTTOCACCA	TGGGTTTTTG	AATCAGGGCC	CCAGATAGAA	CAAAAATGTG	GTTATTCAGT
	CARARAGEA	TCACITGITG	AGAAGACTCT	TTTTTCATTG	AAGTGTTTTG	GCACCCTTAT
49561	CAAAAATCAA	TCTACCATAA	ATGTGAGAGT	TTATTTCTGG	AGTCTCAATT	TTATCCCATT
49621	ATGCTATAAT	CTATAATCCT	ATCTTTTTT	TTTTTTGACA	GAGCCTCACT	CTATTGCCCA
49681	GGTTGGAGTG	CAGTGGCCCA	ATCCCGGCCA	CTGGCTCCTC	CTCCCAGGTT	CAAGCAATTC
49741	TCCTGCCTCA	GCCTCCCAAG	CAGCTGGGAT	TACAGGTACC	TGCCACCATG	CCTGGTTAAT
49801	TTTTGTATTT	TTAGTAGAGA	CGGGGTTTCA	CCATGTTGGT	CAGGCTGGTC	TGGAACTCCT
49861				CCAAAGTGCT		
49921				TCAGGACTAC		
49981				CAACTTCAAA		
50041				TCCCTAGGAA		
50101				TTTTCATAAG		
50161				AATACTGGGG		
50221	AGAAGAAGAA	TAAAAACTTG	TCATATAAAC	AAAAAAGAAA	TGACCAATCA	CATTGTGGAA
50281				GCAGAGAAAT		
50341				TTAGAAGACT		
50401				AATAGATCAT		
50461	TAGGAAGCTA	TAAATCCAAG	ATTAAAAAGT	TGACTGAACT	GTTAAAGAAG	AAACTCTAAT
50521	CTTGAGCCAC	CCTATCCTTG	CTCCACCTTC	TGCTGCAAGC	AAACAGAAAT	GCTGAAATTC
50581	AACACTCACA	AAGGCTGGTA	AGCTGGAAAT	GACAAAAATT	ACTCCTGGGA	AAGTCAGATT
50641	TAGAATTAGG	CCATATTTGT	TGGGGTTCAG	ATTTTCATGT	ACACTTGGGA	AAGGGTTTAG
50701	CTTATAGGCA	CATGCATGAA	GGGAACTGGT	ATAGGGCTGT	GTTCATAAGG	TCAAGAGTTG
50761				TCCCAGCACT		
50821				CCTGGGAAAC		
50881				TGTGGTGGCA		
50941				GGGACATTGA		
51001				ATGAGACCCT		
51061				TACTTTTCTC		
51121				GGGGTGAGAG		
51181				TTTTCCACCC		
51241				AGTACCTGTG		
51301				TAGTGGAATA		
51361				ATGGCCAGGT		
51421				AGCTGCTAGA		
51481				TGGCCCTGAT		
51541				TTGTTCTAAG		
51601				CATTTCTTTT		
51661				AGTGTCAACG		
51721			The state of the s	TAATGAAAGA		
51781				TTCGGGAGGA		
J	THETOTACCI	CHURCIGIA	ATCTCAGCAC	LICGGGAGGA	JOOGGCGGC	MUNICACITO

Figure 9 (Page 16 of 74)

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TTTTAA9A9/	4 ATT00000001	DOAAABBATT	REATCCCTG 1	TTTADAATTA	ATDDTATDTT	19285
TODADADTDE	D TOTBEATOO1	C SOTTEABEE	(ADTOOTOTOO	TOOTATOTAS	TGGGAGGAAC	10285
DDATTAATTI		TTOODTADAT	ATDDTTAADA	TTDAATTAAA	CONTRACTOR	T#185
TODADAATAA	Y TTATOOTTT:) DTATTAADTI	TADDOTTTDA	ATOTTODAAA	CTTTTGTTGA	18085
ATTTDAAADE	ATTTTDAATI	T STOADATTS1	Z ADATAATATD	ATTOAADATT	GGCCACAAGA ?	28027
AATOTOTOAS	ATDABBTBTE	TOTATTTTO	TOAABOBAAA	DOTITION	DOAATADTAD	19645
DADDOTTTTT	TTCCTTTTTT 7	TATTTOTOOT	DODITITAAA	TATATTTT	TACAACAT	10665
ADDITITIADS	AADADDADTO	TATTTAATTE	ATDAADAAAT		ADAAADDTTA	T 7845
TTCASTCTT	SAADTDDDTA	ATTCTTCGAC A	TTCACAGACA	TOASTACTT	TTTTADADAD	18445
AATDTTTTTA	TTTAAATADE	AACATTTCTT	DATATTAAAT	TTTDATTAAT	SANGGRAD	TZLLS
ATTTACATTT	TOATAADTTA	TTTDTDAAAT	ASASSITSSA	TADATTOOTT	TDATTTTATA	19945
TACCAAAGGT	TTTOOTAATA	ATSCATCCTC A	CCACATTCTT	TAAAAATTOOFF	TOTAATTTTO	T0945
DAATETTTT	TTCAAAACAA	DOAAATTODI	TOAAAAOTTT	DOAATTAACA	TTTOTTTAA	T \$ 5 L S
DADADATT	TOTATAAOOO	ATTTDAATTD	DDDATAUTAT	PICHAMANA	DADTTDADTT	
STAATTTTAT	TTTTDTDTT	SSSAAAAASA	ATTTOUTHE	AMBAMCAGO I	ATTOTTTTA	18745
ATATTTTOOA	ATTTOTTTTO	ADATAT"I"I"I	ADAATUUAAAT	WOORDGE TOTOL	AATTTTTADT	77775
TTATTTATAD	ADATOTOA	AAATAATTI	DATTOAAATT	JIWHITHIIN TO	TOTOOTTATA	19845
TTTCACCTTT	ADATDDAATA	DADDATATT	TICTICTAGE	DEFACET	TATATTTTAD	10572
DOTAAADTOA	AAĐAAADDAĐ	A'I'I'I'I'I'DA'I'I	ALTITIATO	AJOALITAJI	TTAAADADTO	27241
ATTOTTOODT	PATTOOTOAT	ATTTTTTTTTTT	4J0AJ1AJA0	TCCTTTTA	TTTGTTTTOO	18172
ADTTDAADAA	TTTTDAATAT	ATOAACCAAA	I RUMMUU I UM	DDDTDTA A AT	ADTTTTDATT	77775
TTDADAAAAT	Setetasare	0.1.9999941.7.1	TAMAMONJATA	GGGGGGGGG	TODDAOTTOT	19072
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DAADDDADTD	ADATAAADAD	TOUTHURITA	PATCANCEACT	SATAASATAA	AAAADDDADT TDADAAADAA	τ9495
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TTTDADTDTD	DIAAAT IADO	TATATATA	9499471949	ASSASTOTSA	Tectorial Tector	18595
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TOAAOOOATA	GCCCAGTAAA	TOADADODIT	TATTĐAĐĐTĐ	DOODTDAADA	TDAAATDADD	T9855
SATTTOOTTO	TOTION SECTION	GAAATGCTAG	TADTTDDTDT	DODITOTITIT	TATTOTOTAD	T0855
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ATTAGATTA	nnnaaaTT9A	ADDTTOTOT	ADDAADTTDT	ADTOATOTOA	TACAGCTCAC	18055

61561	AGAGATGCAA	TATTTAGGGT	TCAACAAGAC	TGAACTTCTG	ACTCCTTTCC	CTACCTCTCC
61621	AGCATGTTAG	ATTCTGGGTC	CTTCATCCTA	ACCCCCTGTT	CATGCCATAG	CCACCCTGTG
61 68 1	GTACCAACTT	TGGAAGCCTG	GATCTTCATC	CCCTCATGAT	AATGAGTGTC	CCATTCAGGT
61741	CTCCATGCTC	AGCTTGGCAA	GAGTATCTGT	CTTCTCCTCA	TGGGACGGTC	ACATTCACCC
61801	AGCACTGACA	GGTTCCATTC	CCACTAGGGT	GGCACCCTAT	ATGGTCTGAG	TCCAGGCCTT
61861	CCTGGTCCCT	CAGTAATCTC	AGCATGGTAG	CACAATCGAA	AAGGGCTAGG	CACGGCAGCA
61921	CCATTTCCCA	CCAAGAGGTC	TGATGGCTCA	TCACATAGAC	TGAAGGAGAT	TCTGAAGAGC
61981	AGAGGTGGAA	TGAAGAATGA	ATCCTGGGCT	CTGCTCTTCC	TAGGCCTGTC	TTCCTCTCTC
62041	CCGAGATGTT	AGCTAACTCA	TGAGAGCCAG	AAACCAACTG	CAGGCTGGCC	TCAGGCACTT
62101	AGGTAGTGCT	TCAGCCTCAG	CAGTCCACAT	TCTAGGAACC	CTCATAATAT	GGGTTGAAGT
62161					CCAGTACTGA	
62221					GCTAATTTCT	
62281	GCTGCTCAAA	CCTCTAGGGG	AACAGTAAGA	CGGGCAGGTT	GTGGGTCTCC	AACCCCATGA
62341					AAGCCACAGT	
62401					GTGTTAACCA	
62461					ATCCCAGGTT	
62521	TGTACCGGAA	TAAATCAGAC	CACACCTGGG	CTTAGAGAAA	GAGTGCAAGG	TTTTATTAAG
62581	TGGAGGTAGC	TCTCAGCAGT	TGGGCAAAGC	CAAAAGTGGA	TGGAGTGGGA	AAGTTTTCCC
62641	TTGGAGTCAG	CCACTCAGTG	GCCCAGGCTC	TCCTGCAACC	ACCCCAGTCA	AATTCCGCCT
62701	CATTTTGCCA	GGCAAACGTT	TGTTGTGTGC	TCTTCTGCCA	GTGTGCTCCC	CTGGACGTCC
62761	AGCTATTCGT	GTCTTGTGGC	AGGCCAGGGG	AGGTCTTGGG	AAATGCAACA	TTTGGGCAGG
62821	ÀAAACAAAAA	TGCCTGTCCT	CACCGTGGTC	CCTGGGCACA	GGCCTGGGGG	TGGAGCCCTA
62881	GCCGGGGACC	ACGCCCTTCC	CTTCCCCACT	TCCATATCAT	TTAAAGGGAC	CATGCCCTTC
62941	CCTTCCCAGC	ACTTTCCCCC	TCCTGTATCA	GGACCTGTGA	ATGTGGCCTT	ATTTGGAAAT
63001	AGGGTCTTTG	CACTTCATCA	GTTAAGATAA	GAGTGGGCTC	TAACCCAACA	TAAAGGGTGT
63061	CCTTATAAAA	AGGAGAAATG	TCATACACAG	AGACTGACAC	CTATAGAGAG	AAAATGTGGT
63121	GAGTAGACAC	AGGGAGAATC	ACCATTCAAG	TCAAGCAATG	AGTCTGGGGA	TACCAGAAGC
63181	TGGGAGAGAA	ACCTGGAACA	GATTATCCCT	CATTGCCTTC	AGAAGGAATC	AAACCTGATG
63241	ATACTTTGAT	TTCAGACTTC	CAGCTTCCAG	GACTGTGTGA	CGATAAATAT	CTGTTGTTAA
63301	GCCAACAAGT	TTGAGGTACT	TTGTTACTGC	AGCCCCAGAA	AACTAATACA	GTAGGTACTA
63361	TGGACTGAAT	TGTGACTCCC	CGTCGCAAAA	TTCATATGTT	GAAACCCTAA	CCCCCAGTGT
63421	GATGGTACTT	GGAGCTGGGG	CGTTTGGGAA	GTCATTATAT	TTAGACAAAC	TCATCAGGAT
63481					CAACAGGCCA	
63541	CTCATGCCTG	TAATCCCAGC	ACTTTGGGAG	GCTGAGGTGG	ATGGATCACC	TGAGGTTGGG
63601	AGTTTGAGAC	CAGCCTGGCC	AACATGGTAA	AACCCCATGT	CTACTAAAAA	TACAAAAATT
63661	GGCCAGGTGT	GGTGGTGCAC	GCTTGTACTC	CCAGCTACTT	GGGAGGCTGA	GGCAGGAGAA
63721	TCCCTTGAAC	CCAGGAGGTG	GAAGTTGCAG	TGAGATCACA	CCACTGTACT	CTAGCCTGGG
63781	TGATAGAGAC	TCCATCTCAA	AAAAAAAAA	AAAAAAAGAC	AATAGAGCCA	GGTGCTGCAG
63841	CTGATGCCTG	TAATTCCAAC	ACTATGAGAG	GCTGAAGCAG	GAGGCTCGCT	TTAGCCCAGG
63901	AGTTCAAGAC	CAGCTTGGAC	AAAATAGTGA	GACCCCCAAC	TTCTAAAAAT	TTAAAAAATG
63961					TGGAGGCTGA	
64021					TCCAGCCTGG	
64081					TTCTCTCCTC	
64141	CTGCCTAAGC	CCTACAAGCA	CAAAAAGGAC	ACCACATGAG	CACATAGTGA	GAATGCTGCT
64201	GCCACCAACA	AGTCAGGAAG	AGAGCGTTCA	CCTAGAAACT	GAATTGGCCA	GCACCTGGAT
64261	CTTGGACTTC	TGAGCTTCCA	GAACTGTGAG	AAAGTTATTT	TTTTTTTAGC	GACTAAGTCT
64321					AGAAGGGATG	
64381					TTAGTCTGAG	
64441	ATGATGAATT	ATTTTTAAGA	ACTTTTAAGG	GATCTGACAA	GTTTGCAAGA	GCTAGAGAAT
64501					ATCTTTCCAC	
64561			= :		GTGACTGGTG	=
64621					CTACATGTAG	
64681					AGGACTCACC	
64741	CTGGTGGCCA	AGCAGCAATG	GCAGGTAGTA	CACACACAAG	AGGCAGATGA	TACAACACAT

Figure 9 (Page 20 of 74)

Eigure 9 (Page 22 of 74)

DDADDTADA 1	DIDIDETET	DADTDDADDA	TOADADOTTO	TODIDITIO	PEANCTGCTG	17771
DOTODITOOI	T TODAAATATA	AATTAƏƏƏTƏ	AAADADDTAA	CAAGCTCATG	GCAGGGTAAC	19114
DTTDAADDA A	S DADADATTTT	ADDDTTATTA	TOADDOTOTT	TATOTOADTO	DTAATTTAAT	τοττι
SCACATTACA) TATOTOAADO	ADDDTAADDD	TADTODIDID	TOATOBBABA	DTAAADTDD	TOOTL
TATAAADDTA	(ATOTABBAAD	DADATADTAT	AACAGAAGGA	ADAAADAAAA	AAAAAAAAA	18607
DDAAADDAD1	T DDDATTOTTT	DTADDATDAD	ADDAATTDAA	CCAGCCCTCC	TADDOTTOTO	12607
TDDTTTADDI	T TOTTODACOA	ADTAADAAAD	CACATAATTA	ASSAASSSAA	TODADDDADA	19804
DTTTCCTTTC	TOODDDATOO	ATDATODDDD	TAGGCTTCTT	TADTATTOTO	TATOTOBACO	10807
TOOTOOTOTO	TTOTOTAOTO	ADADATADAD	ADADADADET	TOOTTAATAT	ADADADETAA	T\$404
ATOOTOAOOA	TDDADADATT	TOTAGATOT	TOTOAOTAAO	ASTBASASTA	DADTDDTADA	T8904
TADDDDTDTA	CAGAACAGTG	DOTOBABTBA	DDAADADDTA	STASSTTSSA	ATASTASSTS	17907
TTAADAAADE) STICADODII	TOTTAADAOT	ADTOTOAOTO	TATTDATDDD	TOADDUTTUA	T9504
AADADATADD	TTODADADAD	STATSSAA AD	TATOOTOOOT	DOADADBAT	DDDTTATATA	T0502
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ATOTAATOTA	CAAAACTTTC	TAADTATADT	TTAAAAATAA	TADAAADDAA	TABILITIONIA	18507
AADATTAATA	SAASTSAAS	TATOTATOTA	ADAAATATAA	AAADTADDAA	TOAASTITSUI	10207
DAATATAAAA	DDADTATDAA	AATƏTTTƏAT	TASTTSSTTT	TTATATATA	PI'DDAAT'DA'I	19707
ATAATDAƏTT	. TTDTTATTAA	TOTOADAA	DOTTBBADDIT	TADABADTDA	ADTTTAADTT	10207
AAAADDDDAA	AATDDTDDAD	CAAAGTCACA	COACTCACT	ATTTƏƏAAAA	SOACASASTO	T7T04
ADAABABTDT	ADATTTDTDT	STTATTATTS	DOATDDADAA	ADTADDDADT	AATADTDDIA	T8004
ATTOOTOAAT	TOTAAATTTA	TOAADAATOT	TOTTODADAO	TOTATAAAAT	LITTERESTEE	12007
ATATATƏTAT	ATAATTTATT	ATTDAAAADA	TTTTDTADDA	Teearurae	AACACACAA	19669
TAADDADDA	DTTAADADDT	TDADAADAAD	ACAACACCAC	ADAATTODAA	AADAAADAJI	10669
ADTBADAATA	DTDATTATAA	TTATOTOTOT	DAATTDADTT	DDDAADDADA	2515512251	T 7869
AATATADOTO	ATAAAADTAA	DACTEAGEDEA	DOTARDATAD	TOTALTOUR	STRACALITAD	
TAADADTDDA	ADDDTTATTD	DTTTADDTDT	TADTOTTAAD	TOTORANGE	TITITITE	1869
DOTOTOAOTA	ATATTTAĐTA	DDTADDATTT	STSTSTSTAT	AATTIJATA	TIPHILIPPI	12769
TTATTCTGTT	TTDAAAACAA	AAAĐATAĐAT	ATCCATTTG	DITAATITIA	TELOGOGIE	19969
ATDAATTDAT	TATATAATD	ATTOATOTO	STASSTTSES	AAAAATTAAA	##ODDDDDTTD	10969
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TAAAAADAD	TODDADTTDD	สออาวววอลอ	TTTOOTAGOA	シントンシントシェン	DATATA ADDA	12769
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TOAADATDDA	TATOTOAATO	AAAAAASAss	AAAASIJAJJ	נפטטענורופו	TAMARAMT	18169
AADTOTODO	TATTAADADT	SOSTSACOST	ורונשטוטט	TOTOTATE AT	4777777	12169
TGTTCTTGCC	ADDDTTDDTT	D'I'I'DADADAA	TOTOTAGEOT	PADPTADDTA	AつのTのTつつのT	T9069
TADBATDBAD	TAAƏTAAƏTƏ	Abbrorrasi	TORICACT	TADARDITO	47777774	10069
ADATBADTDA	ASTSSSSTST	TOTTTIATIO	ADIDIOIDAD	TADAADTTDT	エンシスクン・・・・	17689
ADTTOTTATO	COTABATOTT	Tracorroca	AGEOGRAPHA	ADATORAGE	ANCIOLINA	T8889
DTATADITIT	COTOCOTORT	DTDDTAAAAT	5556755666	DIDOMITTO	なてならなつてつてん	TZ889
ATTOCOTTTO	DATTTTTATA	ATTAADAAAA	CASSOCICE	DATITIADIO	エタエンフスをみなる	T9189
SOOSSOAAAO	DDATTDTTDT	TOADTITOATI	PERFERENCE	DIMITTANTE	ST487848A	10789
DATATOTITO	TADDATD	ADITIAADDO	AUAA JUUA J	つてんつてらてつてら	エムコエアをエンス	17989
AADTAATAAA	TOTATATADT	INITACKING	TOARDADAT	られつつれてつられて	TTAATATTTT	T8589
DTAATADATA	DTTATTOTTA	10177K7K77	ASCETEMENT	DATTATTATT	ADAATAATTA	T2289
ATAAAADTAA	DATTTƏƏTƏT	AADDOADITI	DIDIDIDIDI	ADADBBATTD	TOOTTOOOTT	T9#89
TODIOLOGO	TOATTTOADO	1 J JANIAJAI	PTPTPTPTPT	DOADTITIOTT	STEEDETATT	10489
AADADATAAT	TABADADTTD TBATTTDADD	MILLILLIA.	###74507T73	TOTTATTOTT	nna4aaaaaT	17689
TDATTOATTAA	AATITIDIDAM	AIJJULITE	AAASTAATTT	AADTITITATA	DATTAADADT	18289
DTAAAAADTT	AATTTOTOAA	#####################################	TANTETATAT	TTDDAAAAA	AATTDAAAAT	68221
SAAAD'I-I'ADA	DODATODIT	moroorammo	TTATTOTTAA	DITTTTOTATO	CAAAGCAAAC	19189
DATOTACION	TTGGCTACAG	サントンエムタイン	TTADATOOTA	CACAACAAC	DAAADTOTOT	TOT89
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74501						
74521	CTAGACTGAT	TTAAAATGTT	CTAAAAGTGT	AAAATACACA	CCAGGTTCTG	AAGATTTATC
74581	ATTTAAAAAA	GAATGTCAAC	TGTCTTTTTT	TTTAGCTTAT	TTATTATATG	TTGAAGTGAT
74641	AATAGTTTAG	ATATATTAAG	TTAAATAAAA	TATCTTAAAA	TTAATTTTAC	TTGTTTCTTT
74701	TCATTCTTTC	AATGTGACCA	CTAGAAATCT	GGAAAGTATT	TATGTGATTC	ACATTCTATT
74761	TTACTGTCTA	GTATTGCCTT	ACATCATCAG	GTACCCCATA	AGTAGGCTTT	TTAGATAATT
74821	CTCTAATATA	GCTTGGAAGG	ATATGGAGAA	ATATTTTTGC	GTTGCTTTTA	AGTTTTGCAT
74881	AACTTTTTCA	ACACACTTTA	TAAAGGATCT	AGAAAAGGGT	TGGTTACATG	TTTCTCTGTC
74941	TTCTGGCCTC	CACCATGTTG	CCAGGAGGTT	GGGGACAAGA	TTCTGGGTGG	CTGGATGTCC
75001	TAATGGCTTG	AGGTCTGGAC	TTGAGATTTG	CATATAAAGA	GATGTGATTA	GATTGAGTCG
75061	ACTAGAAAAA	TCATATTAGA	GAACTGAATC	ACAGCGATTA	AATTTACATG	TCGATTTATA
75121	AACCAGGACA	CCAATTTATA	GTGAAAGAAG	GTCCAGTTAC	CTGGTAATCA	AGACGTTTCA
75181	TAGCTATTTT	CATGATGGAT	ATACTTAGCT	GAGTTTTAAA	TGAGAAGGGG	GTTCATTGCA
75241	CATAGAATAA	GATCTAAGTG	AAATGTTTAT	TTATTTTTTT	TTTTTTTTGA	CATGGAGTCT
75301	TGCTCTGTTG	CCCAGGCTGG	AGTGCAATGA	GGCAATCTCG	GCTTCTGGAG	TGCAATGAGG
75361	CAATCTCGGC	TTCTGGAGTG	CAACGAGGCA	ATCTCGGCTC	ACTGCAACCT	CCACCTCCCG
75421	GGTTCAAATG	ATTCTCCTGC	CTCAGTTTCC	TGAGTAGCTG	GGATTAGAGT	TGCCTGCCAC
75481	CACGCCAGGC	TAATTTTTGT	ATTTTTTTA	GTAGAGATGG	GGTTTCACCA	TGCTGGCCAG
75541	GCTGGTCTCG	AACTCCTGAC	CTCAGGCGAT	CTGCCCGCCT	CAGCCTCCCA	AAGTGCTAGG
75601	ATTACAGGCG	TGAGCCACCA	AGCCTGGCCT	AAGTGACATG	TTCTTATATT	GTTCCTTTCT
75661	TTCTTTTTTT	TTCGACTGAG	TCTCACCCTG	TTGCACAGGC	TGGAGTGCAG	TGGCGTCATT
75721		GCAACCTCTG				
75781		CCCCAGCTAA				
75841	CTAGGCTGAT	CTCAAACTCC	TGGCCTCAGG	TGATCCGCCC	CCGAGTCTCC	CAAAGTGCTA
75901		CGTGGGCCAC				
75961		AGGTGCTTCA				
76021		CTGAGGAATA				
76081						GACAATGTTT
76141		GACAAATGTT				
76201		CATTTGTCAT				
76261		TTCTCTTTAA				
76321		TAATATTACC				
76381		TTTACTTTGC				
76441		AAGAAGTAGT				
76501		AAGAGGTGAT				
76561		TTTTCTTGTC				
76621		TCATCTTAAA				
76681		GATATATTTG				
76741		ACTGAAAGTA				
76801		AATAAATAA				
76861		GCCAAATCTA				
76921		ATCTTGAGGG				
76981		AAAGAGCCCA				
77041		AAGAGGGATT				
77101		CACCCGGAC				
77161	TGCTTTGGAC	TGGCCGTGGA	ATCCTTGTCC	CAGTCCACAG	TTCCTGTGCG	ACTGCACGAA
77221	GAATTCACAG	AGGACCTGTG	TTACTTCCCT	TGTGAAGAAA	CAGAATTATC	ATGAAAATTT
77281		CATTTCGCTT				
77341		AACCTTCAGG				
77401		ACTGCCCCAG				
77461		CGTTTGTGGA				
77521		CCAAACCCAA				
77581		CAAACACAAC				
77641		GCACCAATCA				
77701		ATCGGCGCTT				
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Figure 9 (Page 24 of 74)

Figure 9 (Page 26 of 74)

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AAAADTTDTA	A TOTAAAADT	ADDTDDADA/	TGTTCCTCTA 3	T-I-DAAATTOOA	TITCHCHICH !	14958
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AASSSTTTT	N ADAADAADDE) TDADATDDDA	TTTADDDATA	. ADADAADAU	HIMPOHWKIN	19569
A DTADATTA	A DADADAADDI	CAGGCCCAG 1	ADTABABACC	ATADITIONE	TIGOTOTIVA	83287
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つずつつむをつまつご	D STOOTSTIAM	SOAADTTODD	TCCACTTCCC	CACTGCAGCC	TIBBLISTED	18678
TROUBLE) TEASETTEEA	. DDDADTDTDT	 DTDTDTDATA 	DASASTITITA	LIIWIIIWII	87921
TATTTATTT	. TTOTOTITOA	AADDITATIA	SETAATETTS	ATOUGHERT	AAAAATADAA	19878
TOADTATOAA	AATADTDTDT	TOTTOTTADA	DTATITAAAA	DIACOMACT t	HAIRUIJAA	10828
AAATADDTTA	TAADTAADTA	DAAAAADDDA	ATABTSBTSA	DAMAGASSIT	AATAAASTTA	17728
TAATAATƏƏD	AAAAADTTTT	AATAAAATAT	TTTDATTAAA	TITIMOTIT	TTADTATATT	85681
AATDATTADA	DITIAATITA	TATTOAAAAO	TTOTTATADI	DOLKORD LOCATION	DDDTTTAATA	82621
ADATDAAADD	PACTITITATG	TAAAADTDDA	DTTDDSTTAA	COTTOACTO	TODADATOOT	19528
TOTOOADADA	DABAABABAA	ATDDDTTTAA	AbbbAAAbA	STOCKSTEE	TTTAAADDDA	82507
ADATTTTDAD	ADDDDTTDAD	DATATODIO	TTTTTTTTT	DI INNOMATA	STSATSTATA TTTTA 4 4 5 5 5 5 4	85441
DOTTDDAADD	DTDADTDTAD	TDADADDTTT	TTOTITIONAA	DETAINITHMI	AAADATADAA DTDATDTATA	18528
ADTAAAATAD	AADDTADDAA	AAATAADDDD	TADATAAATA	ARJUANA JAUT	DTAADDDTTT 44454T4944	82327
AADDADTADD	DTADTTDADD	TTTĐẠĐĐẠA	ADDAAADAAD	TITALACATI	ATTOADADAD	19228
TOADATTDDO	DDAAAAATTD	DAAADTTAAA	AAAAAAAAAA	DEET LOVOYD	ADDTADATTA ATTDABADAD	82201
TATDATTATT	TTAADDTADD	AAAAAAAAA	DOTADARTIA	SASASSAAAS	ADSTADATTA	82141
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AADTTTTTDA	AATTTTACTT	TTCTTGCCTC	ADAMAMATA	DAATTTTADD	ST442TST3A	12028
	エー・スーン 4 4 D.T. 4	ATDTDDATTD	9913711704	TWOWSTINES	1 7 1 7 1 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7	T96T8
		T427TF3'I'I'I'	WW DIDITIO	TUTUARTER	****	10618
		444453A'I'I'A	WWWATIANA	TUTTTLEFE		81841
		7 4 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7	THAMPOUTTE	TIRILWAN		TBLTB
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		DOTOATORY	77 I WWI DITT	Wawaya		87457
	TOADAAAADT	なつつここれ. リ. プンン∀	THINNWATU	22001222		τ9ετε
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DITIDDADADD	DOADTOADU	TOORGAGGG	シャン シンチャン・	TODTTAADAD DADADTDDDT	DDBADDTDAD	87 7 87
ADDDDTDATD	5 TAATUCCCGG	TOO LOOGUETOUS	つつりつつつるなみ丁	TODITAADAD	ADDDADTDDD	87777
OATOTOTOO	DTAAAbibbi	ボンカイン シン・・・	PATONIANDAA	DODATTAAAA TOOTTAADAD	ATATDAAAAT	19018
		*^ (TCGCC)	20422ABAA2	TTDADDADTD	DADDADTADD	тоотв

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07403	mmanana s s					
87481	TITCTGAATT	TTGTGATGGC	TGTTGTTTTG	TCAGCTTTTA	TAAAATTGGA	ATTTGATTTT
87541	ATTTTCCCAT	TATAAATTTA	TATTTACAGT	CTGCAGTACT	TTTGCATTTT	TAATTTTACA
87601	TTATAGCTTT	TAATAGTTAA	CAAGTTGTAA	AAGGTTTGAT	CCCCAGAAAA	CCTTGATCTA
87661	CCCCCTCAGT	TAAGTATACT	AATATATTA	GAAAATGGAT	GAAATCAGCA	TTTGAATATT
87721	TTTAAATATT	TATTAAAAGA	GGACATGGGT	AAAAGAGCTT	TGCAGTTGCC	ACCCTTCATT
87781	CTCAAATTCC	CTGGATAAGG	ATGACCGCAT	AATCTTTGGA	TGGTCATACG	CAAGTCTTGT
87841	GTATTTGTTA	CATAAATCTA	TTTAGTGGAC	TTTTGGCAGT	GTGTACTGAG	GCCAGTTTCT
87901	TCCACCTGAG	CTCTGACTCC	ACCTCCAGCA	GCCCAAAACC	AATACTGAAT	TTTGGGGTCA
87961	GCTATTGTTT	TTGTGGACTT	AGGTAACTAC	ACACACATTG	TCTTTATGAT	AGCTTTAATA
88021	ATACTGCCAT	CAGAACTAAA	ATTGTCACGT	GGATTAAAAG	GAGTGACGGT	GGTGTCCCCA
88081	GGAGCCTTTC	AATATGTAAG	TATTTACACA	TATACATGCT	AAAAAGACCC	CTAGGAATTT
88141	TTTTAACAAG	GGCAAAACAG	TAACTCAGCT	TGTTTTCTCG	CAGTAAAACC	GGTTGAAAAG
88201	GCCTGATAGA	CTTGTCTGCA	GTTACAAAAC	TTGTGTGTAG	TTATCACCTT	TATATCTCCT
88261	GGAAACTAAC	ATAGACAACC	GAATGGGTTA	CAACTGTTTT	TAAGTGAAAT	TGTGAGTGGC
88321	TCTGAAAAGA	GCCTTTTCAA	TGAGGAAGAA	ACGGGCAGAC	TTATGCCCTT	TCCCCACGGA
88381	TGCGACGTGC	CAGCTGGATA	TCTTTGGGCA	TGATGGTGAC	GCGTTTAGCG	TGAATAGCGC
88441	ACAGATTGGT	GTCTTCGAAG	AGTCCCACCA	GGTAGGCCTC	GCAAGCCTCC	TGCAGCGCCA
88501	TCACCGCAGA	GCTCTGGAAA	CGCAGGTCGG	TTTTGAAGTC	CTGGGCGATT	TCTCGCACCA
88561	GGCGCTGGAA	CGGCAGCTTC	CGGATCAGCA	GCTCGGTGGA	CTTCTGGTAG	CGACGGATTT
88621	CGCGCAAGGC	CACGGTGCCC	GGGCGGTAGC	GATGAGGTTT	CTTCACGCCA	CCGGTGGCCG
88681	GAGCGCTCTT	ACGGGCTGCT	TTAGTAGCAA	GCTGCTTGCG	CGGAGCTTTG	CCGCCGGTAG
88741			GTACGAGCCA			
88801	TGAACTGAGA	GCAAGTGGCC	TTTAAATATA	GTGAGAAACA	TTCTGATTGG	TCCTGTAATA
88861	TTTCAAAAGT	CCCGCGCGAT	AAAATCATTG	GCTGAAGAGT	GACCAGACTG	ATTGGTTCAT
88921			TGAGTTGCCC			
88981	TATCTGCAGC	GACAAATTGT	CTAAAATTCT	AGTTCATCCA	GTCCCAAAGA	ACAGAGTGTA
89041			TTTAAAATGT			
89101			CAGTCTCGCA			
89161			CATGTCGGGA			
89221			GAACAGCAAG			
89281			TTCTCAAAAT			
89341			TTTCTAAAGG			
89401			GGCGGTGGCT			
89461			AGGCCAGGAG			
89521			CAAAAACTAG			
89581			CATGAGAACC			
89641			ACTCCAGCCT			
89701			AAAAATACCC			
89761			AAAATGCCGT			
89821			TTATCGTGGC			
89881	TATTGGACAG	GACGCCTCCC	TGAGCAATAG	TGACGTTGCC	CAGCTGCTTG	TTGACCTCCT
89941	CGTCGTTTCG	GATGGCCAGC	TGCAGGTGGC	GGGGGATGAT	GCTGCGGGTC	TTGTCACGTA
90001	TGGCGCTGCC	CACCAGTTCT	AAGATCTCGG	CGGCCAGGTA	CTGTAAGTAC	ACTGGCGCAC
90061			TAATTGCCCT			
90121	GGGAACTGCA	AGCCCGGTAG	CGACGAACAA	GTTTTTGCTT	TAGCTCCATT	TTCCACGTCC
90181			GCAGCGGAAA			
90241			CAAACTGCAA			
90301			CAATAGAAAA			
90361			TGTTTCTTTT			
90421			TAAATACATG			
90481			TTTCTATTCT			
90541			AAGGGTTCTA			
90601			AGCCGCAAGG			
90661			ACCGGCATCT			
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Figure 9 (Page 28 of 74)

Erdnre 9 (Page 30 of 74)

DTTTTAATT/	A DOAAATADT	TTTOAADAT) STOTOSTA	DADDDDTATA	TATOTTOATA	T 7 T L 6
ADATESATO	DADDIADDA	A ADDADDADAA	TDAAADDDDT	TADABBADBA	CCATAGACAG	T8046
TOATDAADAS	DAAADTDDT	AAATDAAADA	ATDATTTDAA	DOTAAATDAA	CCGCAGTGCA	TZ016
TDADTDDDAT	TTDADAAADA	4 ADDDDDDDTA	6 SOTTOTTO S	CCCCAAGAGA	ADADOTADDD	T 96 9 6
DTDDADAAA E	D DADATTOTAL	r AAAbADDDD1	sererorias	STBATATDAT	DTOTTTDDDA	τ0696
STEETTETS	D DATDDAAAD	DATTDADAAS	DISAABADAT	TOTITOTIO	TTCCTCCTT	T#896
DDDTTTTT7/	(AATADTTTD1	r Telatestei	TTTATTAAAD	ADDAADTTTT	DAADTATAAA	T8 L96
AAAATTAAAA	AAAAAAAAAA	AAAAAAAADI	STDDDTDADA	DOBADADADT	SECCTEGG	TZL96
TOADBOADC	DODOTADADO) DEAETEADET	TODADDODA	DDDDDDAADT	DDDDTAADAD	T 9 9 9 6
DADDDADTDE	DASSSTOAT	r DBADDDTBAT	。 DTつつりつりつす	DOTADTODD	DTDDATTAAT	10996
AAAADATAAA	(AATDATDI	S SOCCECE S	Y ADDADAATTD	DTOOTAOOAD	ADDTADADDA	T \$ 5 9 6
OT DE ABOADT	. Appoppop	Aecoceae	TTTOADDADD	STAATSTSS	DADTDBDTBB	18496
SECCEDITION	TTTAAAAAA	AAAAATTADT	DADTADDAAA	AAAAAT	DADDDADDTA	T Z 7 96
TABABTAABA	SABDBABTDS	SCCGCCACAG (DESTABAADD	DAADDADDTT	GAGGCAGAGG	19896
DDDDDDADTT	DATTAĐĐAĐ Đ	DECTGAGGTG	ADATATOATO	DADOTTDATA	TOOSTATAOS	10896
STSSTSSSS) ADDATTAAAA	AADATDTDTD	GCGAAATCCA	STACAACSAS	TOORYOUNE	17796
SOTTAADADT	. DEASTTDADI	Y ADADDDADTD	ATODABABBB	TTTDADBADD	OTAATOTOO	18196
TACTETETE	ADTDDADDD	DITTTTG	AAAADTTTTA	DDAATATAAT	55JJJJY5Y5 I	17196
DAADADDOOT	• DTATDTABBA	TTTATADTO	AADAATTTDA	TITGAGGCTGC	AASSAATUAT	19096
TADAAAAADD	ACCACATCCC	TDAAAATATT	AATTTTTTD	TTADATDTAA	AAATTOOUT I	10096
SOASSSETSA	ADDIDID	ADADTDDTTD	TADDIDITIES	DADTADADAT	TATTITITIALS	17656
TTTTAATOOO	TOOODADOAD	DESCAAGEGE	DATDABBETT	AATDADDDDT	T.e.c.T.cvecC	18856
COTOTTAADD	AADTTƏƏƏAD	ってこうこうこうてつつ	AADDDDADTD	DDDTDTADDD	TODITAADOLO	12856
ABBTDBBADD	CACTTGTTGC	CGGAGTTTCA	ADADITITIT	Jalalalalalata.	TTTAĐAATOT	19456
TTADATTDAD	TOTATADTAT	TODDDADTIT	DAATTATTAD	AAAATTATƏT	ADDIDBAIA6	10456
ATAAATTTTA	TTACATOTAT	TOOUTTOTAD	LLLCLLLLCC	TADDDDTTDT	TAUTHURAL	17956
TADTTTDDTA	TTATOTTOTT	TADAATAATA	GCACATTTTG	TOADADITTO	TTATTAADAD	T8556
AATT	TTTTTTTTTTA	DTDTAAATTA	ASAASTASAS	DDDTTAAADT	ATTHETCLET.	TZSS6
ToTopoTDAD	AADTTOTODT	CTTCCACAAA	ADTOOTTTTT	TOTOODADAA	CACTCTCACC	19756
47474577750	AASTOOTOOS	TTDAADDATT	DTADATAATT	T.T.T.P.T.T.T.V	TADATATAOT	10156
TADDTADDTA	DTAABBATAB	ADATAAAAA	DDDAATDTDD	TTADAADADA	STITCICIAC	77856
TOADTITITOD	TAAAADTADD	ATTOTACOAA	ATTTAAAAAT	J.T.J.C.P.C.J.J.J.	CHUIDALIA	18256
TTT77757TTA	ATABATBAAB	DDATDTTADT	TTADDTDATT	A'T'T'T'T'T'A'T	STARICIANTS	77756
ADAAAATAAT	TAAATTTT	TATOATOTIT	TOADITITATO	TTABABTBAA	THITITIES	T9156
ATSTATA922	AAADƏTATƏT	DAADDTDTTD	TATTTAAAD	ADATITUTAAD	THETERITE	10156
なつなわれるアナエム	DTATTDTAAT	DTTAATTADA	TACAATAAA	T.C.T.T.T.CCCC	TITIMANAMAN	10136
4449497999	TASTSATTSA	ADTADDAADT	CAGGGGCC 3. PAGAGGGCC 3. PAGAGGGGCC 3. PAGAGGGGCC 3. PAGAGGGGCC 3. PAGAGGGCC 3. PAGAGGCC 3. PAGAGGGCC 3. PAGAGGCC 3. PAGAGGC 3. PAGAGGC 3. PAGAGGCC 3. PAG	D.L.I.I.L.C. 191	21772277	18676
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つつるエエンンサンエ	AADTDTTETA	<i>5542255119</i>	TOOOASTITE	DOADAGAGAT	OFFITTIVE	17676
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	かいかけがなってか	AATTATATTO	AAAATATAAT	PPPT CCATG	WWGGGGTTTWG	18976
	こっていいりつつつび	TTDDADADIT	AAAAATAJJ	CHCHAMMICITY	TITUODIUTO	17976
440400044	エムエムウエダムつエ	PTOATODDIA	TADIADATIA	THITHITE	V2122222	T9546
TATOOTTOOO	AADAAADAAT	TAATDDATAA	TTODISTER	DOTOATABTO	4212222	10506
	ポンペタタタンポンタ	ATOTTOTA	T	WWW T TOWN I W	DITURITOR	10006
TARRESTAR	つかなつかがわびかん	STTDATAADA	ADTAAAbaa'i	WWWSITITE	T T T T T T T T T T T T T T T T T T T	18646
		TTOTTTTTTT		THIRTITIE	VIDVVOITOR	77576
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	ASTUABITAAD	DATODAADAT	STASAAAT 12	DYVIIIOCCIIWY	0111W111W1	10246
エインAAATンAT	TTOTOTOADT	STSAATSSTS	TATCTGGACA	HODDHININD	¥33131.0	T#T#6
B A C A A TOCOCO	エエムムムなつるんん	TTTDTATTAT	TADAADTAAU	WHITTPCWW	VIDVITAVIT	18046
DTTTTAATTA	DDAAATADTD	TTTDAADATO	STOTIONER	DADDDDTATA	パマンムママンスエア	12076
				こくりつうりてるてる	TATOTTOATA	19656

100441	GTTGTTTTTT	TTTTTTTGA	GATGGAATTT	CACTCTTGTT	GTCCAGGCTG	GGGTGCAGTG
100501	GCACAGTCTC	AGCTCACTGC	AACCTCCGCC	TCCTGGGTTC	AAGGGATTCT	CCTGCCTCAG
100561	CCTCTTGAGC	AGCTGGGATT	GCAGCCATGC	GCCACCACAC	CCGGCTAATT	TTTGTATTTT
100621	TAGTAGAGAC	AGGATTCACC	ATGTTGCCCA	GGCTGGTCTC	GAACTCCTGA	CCTCAAGTGA
100681		TCGGCCTACC				
100741		TTTATGGGTG				
100801		TTCATGAAGG				
100861		ACTGACAAGG				
100921		ATGTGGGAGC				
100981		CATTTTTAGA				
101041		AAAAGAGGTT				
101101		ACAGATGGTA				
101161		CTGGGGAAAG				
101221		TGTAAAATTT				
101281		GCAGCCATTT				
101341		TAGACTGGTG				
101401		CTCTCAACAT				
101461		GCTTCCCAGG				
101521	TAGCCAGTCT	GTGGTATTCT	GTTATAGTAT	CACAAAATGG	ACTAAGTAAC	TATATTATGA
101581		TGACTGATCC				
101641	TGTTAGAGGT	TCCTCTACCC	AGTACAAATG	TACTACAAAT	TATATATGTA	TTTTTAAATT
101701		TTCAATAGTA				
101761	à contra de la contra del la contra de la contra de la contra del la contra del la contra del la contra de la contra del la contra de	ATGCATAATT				
101821		ATATTTGTCC				
101881		GGTGAAAGGT				
101941		CAATGGGTTC				
102001		CCCCATCTCT				
102061	CACCTTCCAC	CATGATTGGA	AGTTTCCTGA	GGACTTGCCA	GTAGCAGATG	CCTGCACCAC
102121		CAGCCTGCAC				
102181		GGATTCCCTT				
102241		AATAGCTCAA				
102301		ACTGATTTGG				
102361		GTTTTCGCAA				
102421		TTATTGGGAA				
102481		TCCCTCCCCA				
102541		TTTTGAAGCA				
102601		CGCTTGTTGG				
102661		TGGCTCAGTG				
102721		GAGTTTTTGA				
102781	ACATATCCAA	GGCTCTTTCC	AAAATGGTCT	ACGATTTGTT	TAGGAAGTIA	GAATAGCTGT
102841						CATTTTATCG
102901		CCTTCCTACT				
102961		GATGAAACTT				
103021	TATTTTTCTC	CACAGCACTC	ATCACTTATC	TCTACATTTT	CATTATGTAT	TTACCTTATT
103081	GTGCACCTCC	CACTACAAGA	CAAGTAGCAC	CGTAAGGAAA	CAGGTTGTCT	GCTTTTTCAC
103141	TGCTATGCTC	CCTGCACCTA	GAACACTCTC	TGGCACTTAG	CAGGTTTTCA	GTAAATATAT
103201		TAATGCTGGA				
103261		ATCTTCTTTT				
103321		AACGTGCACT				
103381	GTATAATCTC	TTCAGGGCAC	TATCTGAGAT	AACTTTTTAA	CATCTCCATC	ATGAATCTTG
103441		AAGAAAATGA				
103501		AATTTTGAAA				
103561		TGGGAGAAGG				
103621		AATCAAATCC				
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Figure 9 (Page 32 of 74)

Eigure 9 (Page 34 of 74)

TTDDATTTAC) DAAADTDTAA	TTDAATTTDT	STOODITAAO	ASSSTTTSAT	TTDDTATADT	τοτοτι
DOTOTAATT	COUTTATATAC) TAATAAAATT	NCARACCTTG :	CTATACACAA	TTOABABABO	TFOOTT
DOATADTDAE	DEACAGCAGG (AADADETDTT	AATADADTAD	CAGAGGCATA	DDDADITIDIT	186601
ATADABAAAA	STDDADDATE	TTODTOTODI	AAAAADTDTA	ADDTTDAAAA	TAĐAĐĐAĐAA	109921
AADTOTTT01	TTTDAAATD1	TOTTAADDDA	ADAACTTACA	ADAAAAADTT	CTAAAGGCAC	198601
DADTTTTDTT	: ADDADIADDE	TOTOACCACTGT O	TAAAAADDTA	DDATATABA	COTTETTEC	108601
DDDDDAAAAA	A DABABBTTAE	ADADDAATTA	ATDADDADAT	TATƏTTTTƏT	TATOTAADAA	T\$4601
TTATATTATA	ATTTTADAAA	SOTDATOADT	DITIATOTIT	TADDADTTAT	TTAAAADAAA	189601
STSTTTADTS) TTAAAƏTAƏE	TAADDAATTA	AAAATTTATA	DTTDADADTT	ADTTDTDAAT	109601
STOSTOTTAS	DAADAABADA	ADTODTODO	DDTTTOTTAD	ATTOADTODD	TOADOTTTTA	195601
TODADOTDAD) TODATODIDE	DEDACABETA	DADTADDADT	CATTACTGTC	ATTDAAATAA	105601
DAAAATADDA	TAADDADTDI	DDDTJATDDA	DOTTOTOTAA	ADAAADDATD	TTCTTTATT	109601
TTATATTTT	. DADITJDDDA	TATOTADTOT	ADDTODIA	DATATTDADD	TTTTTTTAT	109381
DAADTDDAAD	TODACCODED	CACAAGCTCT	TOATOADTOD	AADDDTATOT	PETTOADDITA	125601
ADDDTABABA	TTOTATADAT	ATDADADA DA	DOTATTOTIT	ADABATADBB	TOAATOTOAO	197601
DDAADATATT	. DTDADTTDTT	TTTOOTTAOT	ATTTGTTTD	ADTOTATAAO	TAAAASTATT	102601
AAAAATTTTA	, DOTTTATAAT	DAAATATTTD	STTTTT	DAATAATTTT	STGTTCCAGO	191601
TTTADTTAAA	, TTDATTTƏDT	ADDITAADIT	TAAASTABAA	ACACACACAA	DTAAADAADA	180601
ADAADTƏTDA	CGCCTGACCA	TAAAATDATA	DOATOTADTO	AAATƏTAAƏA	TOTADATODO	120601
STICTITICIT	ADAATDDADA	TTTTTTATAA	TTTACATT	TCACATTCTT	SOSTITOSABA	196801
AADDTTAATD	ATDAAAATTA	DAADDTOTAA	ACAACCACAG	ADATTTDAAT	DADADATTTT	106801
ADADTADATT	TTODATATAO	TATTTTTTA	TOTTTTATAT	TTTTAAAATT	TDATTTATA	108841
TOAAAADAAA	DEDATETACE	AASSSTASSS	ADATTTAAAA	DAADDTDATD	ADTIADDDDI	T8480T
TƏƏAƏƏTTIT	TGCTTTTCGG	TOTABTTETO	TTTCTTTAD	DAATTTTDAA	DATITIAADAT	108921
DTAAAATTAT	TOATOTOOT	TOTTOATOAD	LLCLLLLLLL	ACACCACA	PTACOACOTA	199801
TOAGGATOTO	OTOTOTTTO	TTADDTTDDT	DTATDTĐATA	AATOTTOOTT	ATGTGTGGCT	109801
TATTTTAAD	TATATATTOO	DTAATTTAAT	CGACCAAGCA	TOTOTOAOTO	TOTAAATTTTO	195801
ADDDATDDAT	DDATADDTAT	TTTADTDTD	DTDAATDTAD	STGAGGCAGG	TAAAADəTəə	189801
TOTTATTTAA	ADTODAATTA	DAATTDTDD	TTTADTTAAA	TTDTDATATA	TATATATATA	108421
TATATATATA	TATATATATA	TATATTATAT	GGTTTTCTCA	ATDDDDTDAT	ADATAADTO	198801
ADTOTTTTAD	AADDATTODT	TTCCTCAAAA	DTDTDADDTT	DADDDTTOTO	TOTTAAAAA	108301
TTADAAAADT	TTODAAAOOT	TTTTADDDAT	DATTDDDDDT	DAATTDATDA	ATOUTOTAMA	108541
AADDDAATDA	DAATDTTTDA	ADAAATTTDT	TTAGGATTAG	TATOOTTOOA	TOTODON'S T	181801
TADTATDADD	TOAATOTOTO	DAADADAD T	TTATDDTDDA	ATTATTODAD	TOTITATION	108151
ATSTSASSIT	DTDADTT	TOOOAATAAA	DADAATTADT	TDAAATTOTO	DIATOADATI	T9080T
ADDAAADDDD	DDDDDADDAA	TOTTTAAAAAA	TCTTAACAGA	DJ.J.D.J.J.J.J.J.	AATAATATA	T0080T
ATDAAADDAA	CACTTATTCC	CAGAGCCACC	AGCTCTTTT	AADDDAAAAT.	TOTTOTT	T#640T
TTTATTTOATTOA	AATSTAAATS	SAGAAABAB	CGCATTCGCG	GCTCGCTCGC	ANGACATCCA	188401
ADDDDTATTA	TOADTDADDD	AATOOTAOOT	ADDDDDTTTD	SAAASASASS	VOLULT TOTON	128701
PATPOTTDAT	SSEASTETT	TGCAGGAGGC	OBOBBLYBLE	SAGETERE	Y221125211	192201
TTABDDABAA	DITTOADAADD	DEDITARABAD	CCCCCCCCCCC	9Y22119229	LJONNOOJII	TOLLOT
よってつつかてらるら	22422TDAAA	ADDATDBDDB	DOTABA BUBU	101000101	7470007770	T 7940T
2247222242	TOOBAAAAD	Tもつももつももつつ	GCGCCGGCTA	COCAAGAGG	201200002	TBSLOT
440000000000	944792422 9	DEAAATEEDE	DODADOTAAA	PACAGE TOGG	WANNETANTA	103201
つていららてよつつら	TTTTDDTDTD	TTTTOOAAOO	DADDITUTE	OVERVENCE		199201
TODATOADAA	DAAAATATDT	DADTDTDDTD	DTTDATAADD	TELLIGER	TAADTTTTTA	107401
999999999	AADAADDTAA	DOADSTADAS	ACCAGGCACA	ADDITIONAL I	מפשמרנורות	T&E701
ADDDTAATD	DOATODATDA	DADTODADTA	GGTTAACAAA	DDAA TODAOU	ATATABBITAA	102201
うようつずらずずつA	DAADATDADD	TITITOAAADD	ADTTDABAAT	TTTAAATOTT	DOTOTOTITO	107221
STTSDSDDDTT	TOOADOTTAA	DAAABATDAA	TTTTATATTT	TTTOTTTABI	CHATATICTC	191401
STSTESTIAAT	CAPATTTCCC	DTATADDDDA	AATAAATTTT	AADATAAAAA	シ はフラフセフセッコ	
STATTOTTOA	DADDDDTTAA	TOTTTTTADD	ADAATDƏƏDD	วยว∀ววยววย	ADADDDDDADA	T01L01
TITTASSSTTSSA	TDADIDITIO	DADADTTDTD	DTABDBAADT	PAACTCGGGG	OTODAATOTO	T0600T
ADTDDADAAT	AGCGGCGTGA	SOTOADOTSO	AADDDDDTDT		TOOODITITIO	186901
						126901

113401	GCCCCCTCTG	ATGTAAGATT	CTCAGATGAC	TTGCATCTTC	ACTGTACCTG	TCAACCCAAT
113461	AGTCTTCTAT	TCCTGCCTTA	AATTGTAAAT	TCCAAAACTG	ATTTAATTGT	GAAAGTTTCA
113521	AACTGTACGA	CCTAGGAAGT	GTCAAAGTTA	GGTGACCAGA	TTTTTAGAAG	TCAGCCAAAT
113581	ATTCAGCATC	TTTGATTTAG	TAACAAATAT	ATTGATGGCT	ACTTCAGCAA	AAAAAATCAA
113641	CTTTGTTTTC	TGGTTACTTT	GCTAACAAGC	TTCTCCTGAC	AGGAGGATAT	AGTGAATAGG
113701	CAGTTGAATA	AGTGAGTTCG	GGTGAGAGGT	CTGAGCTGGA	GATAAAAATG	TGTGAGTCAT
113761	CAGCAGATAA	ATAAATGCTG	AGACCAGATG	AGATGGCTAA	AAACTGAAAC	ATAATGTAGT
113821	GCAGCATTGT	TTGTAATAGT	AAATGAGTGG	CAACTGTAAA	GTTTTCATCA	GAAAGGACTA
113881	GAGTGATCTA	TACATCCATA	AAATAGAGTA	TTTCTCTACA	CAGCCCTACT	AAAGAATGAG
113941	AAAGCTGTAC	TCCACTACAT	ACTCTGGTGT	ACTCTGGCTC	AGTTCTTGGA	CTCCTCTTTT
114001	CTTGGCTAAC	TCAACTGGCC	TCACCACTTA	CATGCTCTGT	GCTCTGTCAA	ATAGTTTGTT
114061	CAACAGAACA	CCACGGCCTA	GCTGTAAGTG	CCACGTTAAC	TTCTAGCAAT	GCCAAAGCCT
114121	GTGATAGTGG	CAGCTTCGGG	CTGTTTCTCA	TTCCCGGGAT	GCCTAACCAC	CTCTCCAAAT
114181	TCTATCAGTT	TGCTTCCACC	CACTTCAAGC	TTCAGAACGA	AACATAGAGC	TTAAGAAATA
114241	TAGGCCCGGC	AAGGTGGCTC	ACGCCTGTAA	TCCCGGCACT	TTGGAAAGCT	GAGCCTGGTG
114301	GATCACCTGG	GGTCAGGGGT	TCGAGACCAG	CCTGGCCAAT	ATTGTGAAAC	CCCGTCTCTA
114361	CTAAAAAAAA	ATAAAAATTA	GCTGGGCATG	GTTGCGGGCG	ACTGTAATCC	AAGCTACTCG
114421	GGAGGGTGAG	ACAGGAGAAT	AGCTTGAACT	CGGGAGGCAG	AAGTTGCAGT	GAGTTGAGAT
114481	CGCGCTATTA	CACTTAGGCC	TGGGAGACAA	GAGTGAAACT	GTGTCTCTAA	ATABGTGTTT
114541	GCAATTATAA	ACCATCTCCC	TGACCTTAAA	TCTCTAGACT	CATATACAAC	TOCATATTTC
114601	ATGTATCTAA	TTGAATAATG	GGCATCTCGA	ACTTGTCCAA	AATATGTTTA	TACCTARAGE
114661	CCAAGTCTGT	TCTTCCTCTG	ATATTTGTCA	TGTCAATCAA	TAGAACTCCA	TACGIAAACA
114721	AGCTTGGGCC	AGGAATTGTG	CAATATTGTT	TGTCCTGAGC	TTCTTACAAC	TICIICAAGC
114781	TGCAGTCAGC	TCTGTTGAAA	ATCAATCAGA	ATACCTTTCA	TTCTTACAAC	TITCACCCAA
114841	CTAGGAGCAA	GCTGCCATGG	CGGTTTGTCT	GAATGACCAC	ACTENCECEN	AACTCCTCT
114901	TGTTTTCACT	TTTAATCCCC	CTGTCATACA	GTTTTTCTCT	AGIGACCCCA	AACTGGTCTT
114961	CTTTTTGAAG	GTATTATGTC	CACTGTCTGC	TGAAAAGATT	CCACTCCCTT	TCCATCATC
115021	TCATAATAAA	AACCAGCATC	CTTATCATAG	CCTACAAGTA	ACATCACCAA	CONTRACT
115081	TTGCCTGACT	CTCAGGGGTT	TCTCAGGGTG	TAAGACTTAC	AGAIGACCAA	CCATTACAGT
115141	TCCAAGCAAA	CTAGGATGAG	CTCCTCAACC	TACTAGATCT	AGIGCIGAAA	CITAGAAAGT
115201	CCTCATTCTC	TTCGCAGTTC	TTTCTCTTCA	CTGACCTTGC	TOTTTOTO	TACCCTCTGA
115261	CATTTCCAGC	ATCAGCACCT	TTATATATAT	TCTTTCTCCC	TRANSCORR	ATGGACCAAG
115321	TATCTGAATG	GCTCTAGATC	TCATTTCATT	CAAGCCTCTC	CHCAAGGGTC	TTGTCCTGGA
115381	AAGAGACCTC	CCATAATCAT	CCCTTCTAAA	ATAAGCTTTT	CTCAAATACC	AACCTTAAGA
115441	TATATACTTC	ACTATCCTCA	ATACCATA	TATATAACAT	CTGCTCATTT	AGCATATATA
115501	ATGTAATAAT	ACIAICCICA	ATAGCATATA	TATATAACAT	TTCCCCACCT	AGAATTATAT
115561	TOTOTOTOTO	CCARCTCCAR	TATATTTTTT	ATAACTAGAT	ATATTTTATT	TTGTGTTTGT
115621	TATCCCTACC	ACCTTCA ACA	CCCCTCLCCT	GAAGGTAGGG	ACTTTGTTTT	GTCCCAGAAG
115681	TGDADGGDTG	TCTCAATTTT	GGGCTGACGT	TTAACAGGTA	GTTTATGGAG	GTTTGTTGAA
115741	GCTAACACAA	TCNATTCCCC	ATCTCATCA	CTCCAGGCTC	TCCACTAAGC	CCACCAGAAT
115801	TOCACTOR	CTTTTACAAAAA	ATCTCATTCC	TTGACCTGCC	ACTGCCTGAA	GCAATCAGCG
115861	TCTTTCTTCT	CITTAGAAAA	TCTGGGGGAT	AGTCTAGGGG	TTGCAAATTA	AGCAACATTA
115921	CCCTATCCCT	BARCAAGGAC	TGCATGAGTG	TTAGGACTGA	AGAAGGCCCA	AGGTGGTGGT
115981	GGGTATGCCT	AAGATGAGTA	TGACATATCA	GCAATGCTAT	GAACATAGCA	ATGCTATGAA
116041	CATATOCOTA	AAACGTAACA	GGAGCTAGTC	GTGGCTTATT	GTTACAACGA	CTATACCTCC
				CTACATTGAC		
116101				GATTTCAACA		
116161				GACATAAAAT		
116221				ATCCAGAATG		
116281				CTTGTGTAAT		
116341				AATAAAGAGA		
116401				TGGTACCAAA		
116461				TATTTCCCTA		
116521	ATCCCTTTTC					
116581	AGGAACATCC	ТТААААСТА	TGGCTACAAT	GGCTTGACTG	GACAAACCCC	AGGCTTCCAG

Figure 9 (Page 36 of 74)

Eigure 9 (Page 38 of 74)

	DTADDBDDAD	DTDTTTDDTT	DDDTDDATDD	DDATDDADDD	DDDDDAAADT	STSSTSSSST	153061
	TอวAวววอวอ	りてつももてもつもも	TODTDDDADT	DDDADDAADD	DITOOTDIOI	DTDDADDDAT	153001
	DESTRUMENT	ADDTDTDDD	DADDAATTDD	DTATATAAAA	DODITIOATAD	ATGCTTCTGG	155841
	AADAATOOTO	DTDDDTTAAA	DOOTITITADD	ATBAAATDDT	TATAADTƏƏ	TTGCGTTTTG	755887
	ATADDDDTDA	AATTĐĐAAĐA	TATTATTTT	DDDADTTDDD	ADDDTDADDD	DAADTTDDAA	128221
	DDAADTTATD	DOAATADTDA	TTAATDDAAD	DTDABDADTD	TOOOOOTOA	ADDOODATDO	152761
	STTTBBBTSS	ADTĐAAATTT	TDADDTAAAA	TTDADDDADD	TAAADTTAAD	TDDAADDATA	TOLZZT
			TTAADAATDA				755647
	DADDTTDADT	AADATTATAT	TAADTDDDDD	ADAADDTAAA	DDAADDDDDT	AAAƏTAƏAƏA	155281
	DATBATETSE	CAGGAAAAGA	TTTADDDATD	DADDBADAAT	TTTDTDDDDA	ADDDAATDAT	125221
	DTTDTTTDAA	DATTDADTTA	AATDAAAAAD	AADDDADDDD	CCCAAGAAAG	AAADDDTDAD	152461
	AAADDDAADD	ASTSSATSS	DAACCEGCCAAG	AADDDAAAAT	DTDDDAATDD	AAATDDTDDD	122401
	DAAAATDDAT	CCCAAGAAAG	DAAATDTDAA	AADDDAAAAA	AACOTAAGAA	CAAATCCTCC	122341
	DAADAADDDD	DTDDBAAAAA	TODDAAAAAD	DOTOADAADT	DODADAAAAA	Secception	1522281
	DADDDDAAAA	ADTDBAAAAA	TOTADDTDDD	DAADDDAATD	AAAAADATDO	DTDDAAA DTD	122221
	AGCCCGGCGC	ADDAAADDTD	つつてつつてもつもも	AAĐAADAADT	DDAADTTDDT	TeesoToeA	15221
	DDDDATDDDA	AADABADBTD	DTTDDAADD D	AADDAATDDT	DDDADAATTA	DDDDTDDAAT	122101
	TĄDĐDĐĐADA	ADAABAABAB	DTDDADDATD	CGGCCGCAGG	DDTDDDDAA	AAATTOTOĐA	TSOZZT
	DODITOTOT	STEGTGGTG	DADDAATDTD	CTCCTCCTC	DDADDTDDTA	STCAGAGCTG	186121
	てもつつててつつつも	DIDDDDDAAA	AAABAADDTD	CTGCAGCAGC	DDAATODTOO	AAADAAADD	126121
	AADAADĐTD	DATTTDDAAA	DADTOOT	CCGCTTCTGC	SCCGCCCCG	TOOUTOADAA	151651
	ADTOTOTADO	ADTBATTDTD	TOOTOTOOTT	TATTTDADDA	TATTTĐĐAĐT	DOLLOGICO	151801
	ADDOTTTTTT	TOODDOOTTO	DDTDDDATAA	CCACCCTATA	CGGCGCAGTC	ASTAASSASS	T b L T Z T
	DDDADTDADD	ออวออออลอว	DODDDATODO	9999943939	SOASOBSTISS	OAAAəbəəəir	189121
	DADADTDTTD	CCCCAGCTGC	DEBABATODO	SEATASSASS	ASSASSSAS	TTDDDTTDDT	129121
	TTOTOTOTOT	TTTTDDDDDA	ASSCAASSBA	DTAADADTTD	STTTAASATS	SSTSTSASSS	195121
	DDDDTDATTD	TASTOTOOA	Təəsəarədə	CTTTTTGAGA	DTDDADAAAA	AATOOOTOAT	TOSTET
	CCTCTCAACC	DODDODAAAT	ACTGCATTTG	TGCTTGTGGC	Abboroorda	AAATAATOTT	15161
	TAAAAATDTD	DABBDAADBB	TOTOTADAAT	DAAAAATTAA	ADDITIONTA	DATITIATUT	151771
	TTCAAAACAC	ATOAAOĐOOT	DDDTDTTTT	TOADOUTDAA	AATOODDOAO	AATOOODUTO	125121
,	AADADTAĐTT	DADATAATDT	AAAAbADDDT	TADAAAATDT	SACACATCGC	Appeorate	192121
	GACGTGAAAG	TTTOTTOTTA	DTTADDATDT	TCTGGTGTTG	TAADTOTTTD	SSASSTTAAT	102121
•	TATOAOTTTO	ADDITIOADAD	DTTTDDAAAD	TGCAAGTGCA	TOAATAAADD	TOTOADUITIA	151361
•	DADDDTADDT	DAADTOOTTA	AADDTTADDD	DAAAADTADD	TATOTTOTAO	55557777447	180121
:	ATAATTODA	AATOTTODAO	ACATGGAACA	อาวอายาอวอ	TTADTODODT	SSTADIBIAD	120121
	DADTTDADTA	ADAAAATAAD	TTƏTTTATAA	ATACCCATA	TOTTODDAGO	OTITI COOPER	120621
	rttadeaddo	ADDIDDITTD	CAGGGAACCG	Tataaataa	ASSASSST	AATTJAaatu	_
1	TTCACACTA	TOUTTOID	DDADTATTTA	Service	TATCCCCAGG	CATANAMALLI	106071
:	DATTDTTDDD	ACAACTTAAA	ACAGCCACAA	ATTOADOTAA	ADTITIOATAD	TIMITOUTT	170841
·	MEDOTABADT	DTAADDDTTA	DTDTDTDADT	SSATTASSST	DITATTTTT	DAIMALLINO	184021 12021
:	DTDTADDTDA	ADATDTTADD	ATAABADDTA	TTDDDDDDATO	DIDIDIDIDIO	Characteric	
:	CCGCATTACC	ADAATAAADD	CAGCAGGAGC	ADDATDAAAA	STTTATTOAD	TCA CTCTCTC	T990ZT
١	ATTDATTDDA	SGAAAAGCTC	DDAAAAADAA	PCTTTTTDA	DDDLLDDLLL	ATTACHTORY	750601 750541
	DADDDTDTAT	TDATAADATD	CAGAAGATGG	PECACTGGTG	AbbATbbAbb	DIAMMATMA	
j	CATGCCTTCT	DTAADTTTAT	AAADTDDATT	AAADATTTTT	AAADDITITIA	TTOOORY	120021
j	TDADATDTA	TCAAACCAAG	DOODATOADT	DAATADTTTD	TOTOORINIO	TITIWYDDII	120021
5	TACAAAAT	TOATTTĐĐĐA	AADADTTTOT	ATATTTOOAA	ADAAT DADAT	THI LOCATOR	150361
,	いなるタンエイエイ	TTDDDTTDAA	AADTTTTDDT	TTATDADDAT	TAATADADTA	WOLDOLDWW	100021
-	STOAAATTT	ATTADDDAAD	ADDDADDAA	AAAADATAAD	CAGTTTAACC	DAAAT TEAAAA	150541
,	としていることと	DEADSTADST	TTADTTDDTA	CAGATCTATC	DAAADA'I'I'I'A	CONTRACTOR	120181
=	TTTTATDDD	ATTTƏƏTTAƏ	TTTTTTTT	TACACCEGAC	ADADAADADI	TAAATIIII	120121
-	ASTSTTASS	TOAAAADTOA	DAAAADTDTA	CATTTTGAAA	TAADDI'AAT'	DADARJULI I	150021
٦	DDATTTAAT	GCAAGGAACA	DTTDDTDATA	DDTDAAADAU	AASSSSSTTS	ADDITUDAROUS	150001
C	DADDDTTD	AGTTGTTCCT	DDDTDDADDA	SOATSSASSS	CAGCACATO	TOURSE PROBLE	118841
			· ·				T886TT

756/162

126361	ACTCCAGAAC	ATTAGGTTTC	AATAGATTCA	TCTGTGTTGC	TGTGTATAAC	· TTTT A A TTT A A
126421		AIGIAATATT	CCATGTTATO	AGTGCDDCDD	TTTNCCTCCC	
126481	CAINITIG	. IICCCTTTT[CAGCTAATAT	' AAACAATACC	CTC B B T B TTC	
126541	GTCTTGGTAT	ATATAGGAAT	ACATATTTTG	TTTGTATACC	TACCACACA	CTGTGTATGT
126601	CAAATGCTAA	ACTCTTTTTG	AAAGTGGTGA	TATTAGGTTT	ACATCCCA TO	ATTGTTGGGT
126661	TAAAACCACA	GTTATAAACA	GCATGGATGA	ACCTCACAAA	CCTAATGCGATG	AAATGAAAAT
126721	GCTGGGAATT	CCTGTTCTTC	CATATACTTC	CCAATATTTT	TTTCCARTGTTG	ATGGAATCTA
126781	TCTTTTGAAG	ATGTTATCCA	TTGTGGCAGA	TGTGCAGTAT	TATOMORMA	AAATTGTTAA
126841	TTACATCTTT	TGCCCATTTT	TTCTTAATTC	GATTGTATAT	CACTGGATTA	TGGTTTTATT
126901	AACAAAAATA	CTAGACTAGG	TAGCTTGAAC	AAAAGGAATT	CAGTCGACTT	GGGCTGCCAT
126961	GGCCAGGCCA	GAAATCCTAA	ATTGAGGTGC	CAAGAGATTC	ACTIMENT	CAGTTCTAAA
127021	TTATTGACCT	GAAGATAGTT	GCTGTCTTAG	ATTGTTTGGT	AGTITUTAGT	GAGGGCTCTC
127081	CCAAATAATT	TATAAAGAAT	ACAGATTTAT	TTCTTACAAT	GCIGAACAGA	ATACCAGAGA
127141	TGGTCGAGGG	GCCCACCTCT	GGCAAGGGCC	TTCTTACAAT	TCTGGTGGCT	ATAAAGCCTA
127201	ATCTCATATT	CAAACCACAG	CAGTCGCCTT	TTGTGTCCTC	TATGGCAGAT	GTGAGATGTC
127261	CATAAAATGA	CCTCATGTCT	CTTCCTTTTC	TTATAAGGAC	ATGTGGCCTC	TTCATATGCC
127321	GGCCTACTCT	TATGACCTCA	TTTAACCTTA	AATATCTCCA	ACCAGATCTA	TCAGACTACT
127381	CTCCAAATAT	AGGCACATTG	GGTGTTAGAG	TTTCAACATC	TAAAGTCCCA	AAATCCCTAT
127441	TAGGCCAAAA	AGATTGTGTT	AGIGITACAC	GGTTTAAGAT	AATTTTGGGG	GAACACAATT
127501	TGTCCTTTCT	TTTTTTTTCA	GGTGGACTCT	TCCTCTCTCT	AGCTGTCTTT	TTGTCCTTTT
127561	CGCTGTCTCA	GCTCACTGCA	ACCTCCACCT	TGCTGTGTCA	CCCGGGTTGG	AGTGCAGTGG
127621	GTAGCTGGGA	CTACAGGTGC	ATACCACCO	CCTGGGTTCA	AGAAATTCTC	CTCCTCCCAA
127681	ACGGGGTTTC	ACCATGTTGG	CCACCACCGC	GCCCTGCTAA	TITTTGTATT	TTTGATAGAG
127741	GCCTCGGCCT	CCCAAAATGC	TCACATTACA	CTCAAACTCC	TGACCTCAGG	TGATCCACCT
127801	TCTGTTTTAA	GTTTTTAAAT	TTTCCTCACC	GGTGTGAGCC	ACCAAACCTG	GCCTGTCTTT
127861	ATTTCCTCTG	TAACTTGTCT	TCACTCTCTC	AACCCTTTAT	CCATTTTATG	TGTTGCAGGT
127921	CTCACTGCAG	CCTCCACCTC	CCACCATCAA	AGAGGCTGGA	GTGCAGTGGC	ACAATCACAG
127981	GTGGGACTAC	ATGTGCAGGC	CACCATCCC	GCGATCCTCC	CATCTTATCC	TCCTTAGTAG
128041	GTGCTGTTGC	CCAAGTTGGT	CACCAIGCCC	AGCTAATCTT	TGTATTTTT	TGTAGAGATG
128101	CCCAAAGTGT	TGGGACTAGA	CTCAAACICC	TGAGCTCAAG	CAATCCATCA	ACCTTGGCCT
128161	ATGCATTAAA	GTCATTAATT	TACTOTACTO	ACCACTGCAC	CCAGCCAATG	ATATCTCATG
128221	CCTTTTTTCT	מדכתדותתון	AAAAATCATT	AAATTAAGCA	CACTGCCCTT	TTATGCACAA
128281	TATAATACCT	TOTTGTGAAA	TTACTTCTCA	TTCTATTTCA	AGGTCATGAA	GATCTTATTT
128341	AGTTGGGAGG	TCTGTGGTTC	CCAATCAACC	AGACTACCCT	CACTTCTAAC	ACCAATTATA
128401	AGAACTTGCT	GAAGCTGTTA	CCARICAACC	TTAGGTTAGT	AATTTGCTAA	AAGGACTCAC
128461	ATGTCATTCC	AATGCAATGT	D D D D TTD TTD	TACAATTTAT	TATAGGATAT	ATAGCTTATT
128521	CCAACAATT	TCACTCTCAC	CTAMBORRA	AACTACTTTT	AAAAAGATTT	TAGCATTTGA
128581	CACATACACA	GCA A A ATTCA	TTOTTTOTT	AGCAGATATG	TGTGCACATA	TATACCAAGA
128641	AAAGATTAAA	ATCACCTCAC	ANANCANA	TAGTTGAAAA	GGGGAAACAA	CTCAAGGAAT
128701	GCAGATCTCC	CTTGCCCCCA	CARCATATOR	CACAAGGCAG	TATTATGGAT	CGAATTGTAT
128761	TGTGGCCTTA	TTTGGAAATA	GRAGATATGT	TTAAAGTCCC	AACTCCCAGT	ACCTCAGAAT
128821	ACAGTATGAT	GGGCTGGTGA	CTTACAACAA	AGATATAATT	AGTTAAGATG	AGGTTATAGT
128881	GTATTCTTCT	AAGGTGGTCA	CCTCAACACA	GTAGTATATA	TATATTTTT	AATAGAACTA
128941	AGCTGCAGGT	CAAGGAATGT	CA A A CCTTCC	GACACACACA	GGCAGAGACT	GCGGTTATGC
129001	GGAAGGATTT	TCCTACAGGC	TTCACTCCAA	CAGCAAGTAC	GAGAAGCTAG	GAAGAGTCAA
129061	TTTCTAGCTT	CCAGAACTAC	AACACAATATA	GCATAGATCT	AATGATACCT	TCATGTCAGA
129121	GCTCTTTGTT	ACAGCAGCCC	TACCARACAR	ATTTGTTGTT	TTAAGCCACC	CTAGCTTCTA
129181	ATGAGCTTCC	AGTTGTCCTC	TCCCACTA	ATATAGGCAC	AATCCAGGCA	AGTTCCAAAT
129241	TGTGACAATA	CACATCACCTC	ACACACASCA	ATGAACAGTA	TTACTTTCCC	AGCATTAATG
129301	GACCTCCACT	CALAIGACGI	ACAGAGCAGT	CCCCACTTAT	GCACAAAACA	TATGTTCCAG
129361	TATACAGACA	CACCUPACION	AACCAIGGAT	AGTACTGAAC	CTATATAGC	TGTTTTTTCC
129421	ACAATAAATT	AGAATATORI	MAGGC I'I AAT	TTATAAATTA	GGCACAGTAA	GAGATTAATA
129481	CTGAAATTTA	CCCTTTTTTTT	TTTTTCCACT	ACTGTATAAA	AGTTAGGTGA	ATGTTTATTT
129541	GAAACCGTAT	DCDDGDGD AC	TOTATTTO	GCAGTAGACC	ACAGGAACTA	AAACCATGTA
		ALAMONUMAC	IGIALITUAC	CCGAGCCTCA (JIGTGCAGTT '	TTAATGGCCT

Figure 9 (Pag 40 of 74)

Figure 9 (Page 42 of 74)

				DADATDDADD		136021
				TOTOTOTTAO		196581
				TOTOTADIAO		TOESET
				DTABBADTTB		132841
				DADAAADAAA		187251
				TOTOCOADAA		135721
DDTDDDADDA	DADTTDADDA	DDADDADDDT	ADTTTADTAD	DADDDTDDAD	TODDADDDTT	199521
TODODATOOT	AATOTOOAOA	DTDDATDDTD	TODOTTTACT	AABAAABAAA	ASTITATOTO	TOSSET
TOAADTODTO	DTTAAAATTT	DTTDATDADT	TOADATDADD	DADOTTOTAD	DTTADDDDDA	732247
DTDTTAADAA	ももすつつもすてつも	ADADDTDADT	DADDATADDT	DOTOTION	TODODIAATT	132481
DDDDAATTTA	ATDADTTT	DTTATTAAA D	ATTTATADTT	DTAAADTOOT	DOTTDATTOT	73245T
ADDBABBAAB	AADOTOTTAT	TOAAOODDOO	DAADADATDA	ATADTOTDAT	ADTOTADODO	192561
TOOTTTOTOD	DDDADADADD	DODADTOTET	COTOBODOT	၁៦၁៦៦៦೯೩೦೨	ADATTOODAA	TOESET
ADADTADTDD	TTTADDDDAA	ADDTADAAAD	ATODAADTOO	TOADDATTDA	TDADTOTAAT	132541
AAADDAADTT	ADADOTATAA	TCTTGTCTTG	TOTOAAOTAO	DADAADDDTT	DDDDDTDTAA	τετςετ
ATTTADTDTA	TADOTTTOTA	TDADTDTAAD	TOADOTODDA	DTDAATDTAA	TATOBBTOTA	TZTSET
DTDAAADDDA	TTTAADTATD	DADTTTADDA	TOAADADDTA	TAAAAADDAD	AATDAATDAA	T90SET
DOTATDADTD	AADDDTDTDD	ATTATTTDAA	AATDADADDA	ADADDTTTTA	TTADDBDTAT	τοοςετ
ATAAATOOTT	DTTTTAAA AT	TADTADTAAD	TTTTDADTTTA	ATDDAAATTD	TADADTTATA	176751
DEADTODOT	DTATCOTED	DTDTABABAD	AADDAADDAT	AADDDTDAAD	AAADDAAADD	T34887
DADDATDDTA	DDAADDTAAT	DDDTAATDAA	DITTIBADA	AGACACAGAA	DOTTDADDDA	134821
ADDIADIADO	ADAAAADTAA	TAADTOTOTO	TOTOTOTODA	DOTAATADDD	ATTTĐAAAAA	192757
TAGCCACAGA	DIDITIDIDAA	DESCATGETC	DTTDDTADDA	ADTODTADTO	ADTITITIT	TOLVET
DTDDATTDDA	ADTTAAAAAA	DTADTTATAT	TTAAATATAÐ	DADATCTGTG	TCAGTCAGGT	134641
GCAATGAAGA	DDTDDTDADD	ATOTTTOAAA	DODITOTAKOO	GGAAACAGAA	TOAAATOTAT	134581
DADTODITITA	TDAATAATTO	DTDTDTDAAA	DADAATDDDA	ATTOACATOT	ATAADTTOTO	134251
DDTDADDADT	TACTOSTOTO	овевссувно	DTADDDTDDA	DOTADDATAA	ADDDTDDADA	198881
AADTOTOTAA	DADDADADAD	TTTDDAAADT	DDDDDDADATA	TDDDDATATT	Deterated	134401
GAGCAGCTGA	DDADTDADDD	DODAADTADD	CTTATTTC	TODOTADOAO	CAGTTTTA	134341
DTDADDDADT	DTDTDDATAT	AAATDDTADA	AAADAAATTA	AAAATAAAAA	TTACCTET	134281
DAADATATAT	AAATADDDDA	TOATADADTA	TAAAADTATƏ	TADATATOTA	ADADATTADT	134551
ADTITADID	TTAATOTAƏT	GACAAAAAC	CCAPAPAPAP	TTATCTTTA	ADAAAATATD	191751
GACCAAAGA	CATTTTCAG	DTDDTDADBD	TATAATOOTA	ADDITOTOTT	TTTDAATTTD	101751
DTTTTTTTTTT	ATADTTDDDD	TATATTADTT	CONTATABAT	STTTSAASAS	TOTOODATT	134041
				DTDATDADTA		186551
				STTATSSSSA		136551
				ATOTOODAO		198551
				DDDAADAADT		TOBEET
				DTABAABBTA		197551
				TOADOADETO		189661
				AGAACAAGA		129661
				AAADAAĐĐTĐ		TOSEET
				GAACAAAGAA		TOSEET
				ATTTAĐADTT		193661
				ADDITOTION		133381
					DOTADATOTA	
					DADDDDTDDA	193561
				TOADTOTOTO		133561
				TOOTAADAAD		193361
					TDAADAAAAA	133081
				ADATAAATDT		120551
				DDDTTTADAD		135051
				AATDTDDADD		135561
				ABBABADDBA		
				1001010000	SSER ATTARAT	132841

139321	AAATGTTACT	CAAAAAAAT	CAGAGGACAT	ATGTGGATAG	ATA ATCCA NO	
139381	COLNOCITO	4 MOGGIIGGG	TGCCCCTCC	L CACCTGTGGG	TCTTTCTCCT	
139441	OVOVOVCIIC	3 GAAAAGAAAG	i AGACACAGAC	ACAAAGTATA	CACAAACAAA	
139501	CAGGGGACCO	GTGTTCAGCA	TACGGAGGAT	CCCACCGGCC	TCTGACTTCC	AAAAGGGGTC
139561	ATTGATCATT	ATTGGGTGTT	TCTCGGAGAG	GGGGATGTGG	CAGGGTCNNN	CITAGTATTT
139621	GGAGAGAAGG	TCAGCAGGTA	AACACGTGAA	CAAAGGTCTC	TGCATCATA	GGATAATAGT
139681	GAATTAAGTO	CTGTGCTTTA	GATATGCATA	CACATABACA	TOTALA	ACAAGGTAAA
139741	GTATTGCTGC	CAGCATGTCC	CACCTCCAGO	COTABGGGAG	TTTTCCCCC	TIGAAGAGCA
139801	TGGAATATAC	AATCGGGTTT	TACACTGAGA	CATTCCATTC	CCCACCCCTA	TCTCAGTAGA
139861	AGATGCCTTC	CTCTTGTCTC	AACTGCAAAG	AGGCGTTCCT	TCCTCTTTT	AGCAGGAGAC
139921	TCAGCACAGA	CCCTTTACGG	GTGTCGGGCT	GGGGGACGGT	CACCICITITA	CTAATCCTCC
139981	AGGCCACATT	TCAGACTATC	ACATGGGGAG	AAACCTTGGA	CANTACCTCC	CCTTCCCACG
140041	CAGAGGTCCC	TGTGGCCTTC	CTCAGTGTTT	TGTGTCCCTG	AGTACTTCAC	ATTICCTAGG
140101	GGAGATGACT	CTTAACGAGC	ATGCTGCCTT	CAAGCATTTC	TTTARCARA	ATTAGGGAGT
140161	ACAGCCCTTA	ATCCATTTAA	CCCTGAGTTG	ACACAGCATA	TETETENCE	CACATCITGC
140221	TGGGGCTAGG	GTTAGATTAA	CAGCATCTCA	AGGCAGAAGA	ATTTTTTTTTT	AGCACAGGGT
140281	AAATGGAGTC	TCCTATGTCT	ACTTCTTCT	ACACAGAGACAC	ATTITICITA	GTACAGAACA
140341	CTCTTTTCCC	CACAGGAGGT	GATGGCCGGA	AGAACATGGC	AGIAACAATG	TGATCTCTCT
140401	ATTGGGAACA	AGCTCTGTTT	AAAAGGAGAC	TTGTGAACAC	CARACACTA	ACAAAACAGC
140461	CTTACAACTG	AAGCCCATGG	AAGACAAATG	TGTACTGCGT	CAAAGAGIAG	AAAGGGTTCT
140521	TAGTGGGACC	TAGGGCACAC	CAGAGAGCAT	בסותכוטכטו	DAGITITAAG	GCAATAGGAG
140581	TCTGCTGGAC	ACAGTGGCTC	ACACCTTAAT	CCTACAACTT	TGGGAGGGGG	AAACATTATA
140641	GTGTAGCTTG	AGCCCAGGAG	TTCGAGACCA	ACCTGGGCAA	CATGGGAAAA	AGGCGGGCGG
140701	ACAAAACAAA	CAAACAAAA	ACAAAATTAG	CCAGGCACGG	TGATGCGTAG	CTCTCCCT
140761	AGCTACTCAG	AGGCTGAGGT	GGGAGGATCG	CTTGAGCCCC	GGGAGGTTAA	CTGTGGTCCC
140821	AGCCATGATA	ATGCCACTGC	ATCTCAGCCT	GGGCAACAGA	CCCACAACC	GGCTGCAGTG
140881	AAAAACAAAA	ACACACCATA	CCCAACCACA	ATGCATCTGT	CTTTA ACTACC	GTCTCAAAAC
140941	CCCTCTACTC	ACTACTAAAT	AGGTGAGTTC	CCAATCCCTG	CTIAAGIACC	AGTACCACAC
141001	ATATTAAAGG	TCTTAGGCTA	GTGACTCATT	CACTCATTAA	ACAAATACTT	TAAGCATGTT
141061	TACTATAAAC	TAAGTACTGT	GCTAGGTACA	AAAGCAAATA	ATCTARGCTC	TATABACTO
141121	ACTTTCTTCA	TCAACAAAAT	GGAGATGTTT	TAGGCATCTA	CTCATCATTC	TCACCTCCAT
141181	CTTTTGTGAC	TGTAGTTGGC	AGAGCTTTTT	ATCAGTTTCT	CTABATAGCT	CTACCACTCC
141241	CTGGTGGATG	CTGGCATGCC	CAAAGGATCC	ATCCTGATGG	CCCTGTCTGC	TTACCAGICC
141301	TGCCTGCCTT	TGCAGCACCG	CTCTGCTCTT	CTGCAGGACT	TCCCTTATCC	TTTCCCCCTCT
141361	TGCTGCTCTT	AGGCTGCTCT	GCTTGTTTTG	ATCTGCTTTG	CATCACATCT	ATCTARACCT
141421	CCTTTCCTTA	TTTACCCATG	ACCAAGGTAT	TATGAGATTC	TGGAATTTCC	CCARACCACA
141481	TTGATTGCTG	GGAGAATAGA	AGAAGTGGAT	TACAAGTGGA	ACTTAGAAGG	CCAAACCACA
141541	AGAAGACGTC	TCTGCAAATC	CATTTAGAGA	GACCTTTCTC	CAGTGGTGAC	TCAAACATCC
141601	AGCTCCTTTC	ATCCTGTGGC	TTGGCCATCT	TCAGCACATG	GCTCCCAAGG	ATGTCCTCAG
141661	GATGGTCTCT	AATCCAAGGA	GCCTGAAGAG	AAAAAAAGGC	ATGGAGTATT	GTGAGTGGTA
141721	GGTGGTTATG	GACCAGTTAT	GGAAGAATAC	ACATCACTTT	TGCCCACCTT	CTACTAACCA
141781	GAACTCACAC	AGCCATAGAC	ACTGACAAGT	AGGACTTAAC	AAGAATCTAA	ТТТТСАСТСТ
141841	AGGAATACGA	CTGTAGCAAA	TATTTAACAG	CTTCAAACAC	AGGTGCATTG	CTATCACTAT
141901	GCTTGGCCCA	GGCCTGTCTC	CCTTTCCTGC	CATGTCACAG	GGGCCAGCAT	TTATCTCTAC
141961	ATTGGGTTGG	TTGGGATATT	AAGACAATAA	TGAACCAATA	CAACATCTTG	AGCATAAAAC
142021	CAACTGATAC	AATGATGTAC	AAGTCAGATG	ATTCTGATGA	TTATGAATTA	TGTCAATAA
142081	AGAAATGTGA	TAACTAAGGT	AATTTTTGTT	TTGGCAAATT '	TTTGTTTGTT	CATGACAGGA
142141	TGAAATCCTG	TCATTTGTAG	CAACATGGAT	GGAATTGCAG	GATACTACAT	TAAGTGAAAT
142201	AAGCCAGAAA	CAGAAAGTTA	AACACCACAT	GTTCTCACTT .	ATATGCAGAA	GCTAGCTAAC
142261	TAAGTAAATA	AGTTTATCTC	ATTGAAGTAA	AAAGTACAAC	AGAGATTACT .	AGAGGCTGGG
142321	AATGGTAGGG	GAAAGAGATG	ATAAAGAGAG	ATTCATTAAA .	ATAAGTTACA	GCTAGATAAG
142381	AGCAATCAGT	TCTAGTGTTC	TATTTGTACT	ACAGAATGGC .	AATAGTTAAC	AGTAATAAAT
142441	AATTTCAAAG	AGCTAGAAAA	GAGGACATTG	AATGTTTCCA A	ACACAAAGAA	ATGAGAAATG
142501	CTTGAAATAA	TGGATATTCT	AATTAATTAC	CCTGATCTGA '	CACTATACA	CAGTATGTAT

Figure 9 (Page 44 of 74)

Figure 9 (Page 46 of 74)

				DATODAAADA		180671
				TAĐAĐĐATĐĐ		149021
				DTDTAADTTA		19687
DATDTATADD	TAADADDDTT	TOTOTITIOTA	AATDAADTTD	ADATTDATAA	OTTODITITI	1068 ≯ T
ADABBBBBT	DETATOTETA	DDTDDTDDTD	TOATOATOTA	AATAATƏTTA	ADTTDAATDT	148841
AADTƏTƏDAƏ	TAATADTTDD	TTTCAGACAT	TAAATƏTAAT	DDDAAATAA	ACAACACCAA	T8781
DDAAADATAD	DOAAAADTƏT	TTADATATTA	DADTADATAA	DTDDTAATAT	DATADTDTOT	1278#1
TOAAADTDAO	DDDATTTTTA	DDADADADA	DDDAAAADTT	DTDAADDADA	DTDDDTTAAA	₹998 ₽ ₹
DATADITDAT	TABAAAADTD	ADADADDTAT	DAATDADABA	AAADATOTOA	DADTATOTDA	T0987T
ADADITODID	STTSSTSSTS	AOTTĐAĐĐOO	TOOSTOOTS	AAATDAĐĐ DD	TODITAAOT	T#88#T
TOTOATODDO	TTTDTDDDTD	DIDDAATITD	DTDDADADDT	ATTOTOTOO	りりてりりてつりつり	T808T
DTADADTDAA	AAABDAADBB	TADDOTTOTT	DTDDTDTAAT	ADDTDAADAA	ADDADADATD	148451
DATDADDOTT	DAATDDTDAD	AAAAADATDT	TTAAADADDT	ATTAĐĐĐĐĐA	ATTOTACOTO	T9E8#T
AATTDATTT	DDTDDATATA	ATDADDTDTT	DADDDDDADT	DDADDATDAD	ADDDDTDTAA	74830T
TOAATADDDA	ATODDDAODA	CTCCTTGCTG	DTDATDDADA	ADADDADAAD	ATDTDADDAD	148241
DAADTTATTT	TDAADDATTD	TOOTOOTOAA	DTDDDAADTT	CCGGGAAGAA	DDATTDDDDA	148181
TTAAAAADDD	AADDAADDD	ADDTADTDAA	DDTDADTATA	TOATADDTDA	DAAAADT DDA	148151
ADDBAAAAAT	DATTDTATDT	AAATTDAAA 9	DAADTADADA	AAA DTADDDA	ADTABDAAA B	T9087T
AAAADAAAAA	DAAAAADAAA	AAADAAAADA	AADAAAAAA	AAAAAAAAAA	AAAAAAAA AA	T48001
AAAADTDT	DTDDTDADAA	DDADADAADD	DDTJJDAJTT	DADETDADDE	TDADDDADTD	T\$61\$T
ADDITEDADD	TAĐAĐĐĐTOO	AAƏTTTƏƏTA	ADADTATADA	DTDDDADDDD	TOATOBADDD	T88L\$T
TAATOTOODA	DTDDDDTDAT	ADDDDDDAT	TAADDDDAAA	ADDDTADAAD	TOOTOODAGO	T Z 8 L T T
ADADTTTDAD	DADDDDADTT	DETTABBTEE	DIDAADDDD	ADDDTTTDAD	TTADSTAAAS	T9 <i>LL</i> \$T
AAAAATTDA D	ATTTATTDTA	ADDTAATAAD	TOTTOATOOT	DTOAATADDT	DAATDDATDA	TOLLDT
DTTTTDTAD	AADAAADƏTT	DAAAADDTAT	TTAATADTTA	TAGCAGCTTT	ATTTDTADAT	T#9L#T
ADADDTDDAA	TCCACACAA	DTATTDAAA A	DTTAADDAAA	DOTATTTATO	ACACTCCTTA	T8547T
DTAADDADTT	ADAATADDAT	TTTOATADDA	ATDAAAAAAT	TOTTTƏƏTƏƏ	AAGACAGTTT	T Z S L D T
DDTTTDADTA	ADSTSTAAAA	DOTAADADTD	DTDATTADTT	DTTDAATDAT	AADDATDTAT	T97L7T
ADAAADDTTD	AGCACCAAA G	SAACACTGTC	CCAAAATTTA	ADTAADATDA	TODATADOTO	10077
ADDATABATE	AADAAAATTA	AADDTDAADD	DATTADTDTA	DIADADITOD	TADAAAADTA	ፒቅፎሬቅፒ
TADSTATSAA	ACACAGATGC	CAAAGAAGAT	DADTTOTOTA	TGACCTGAAC	AGCAGGCCAA	18217
AATTTADT	AATAAAADAA	ACTCAACAGT	AAATTDATAA	DAAADATATA	ADATTTTAT	147221
				DITADADADD		1914
				ADDIDDDITT		τοτέντ
				AATOTOAAAA		τνοίντ
				DAAAAAAAA		186971
				ADADOTTDAD		176971
				TOODIATIOA		T989#T
				DAAADTDAAA		T0897T
				ATTEASTTTA		T\$49\$T
ATADTDDADA	ADAATATTTA	TADDAAAA	AAAAATTTA	TTTTTAAATT	DTDTAAAATA	189971
TTGACCAGTT	TAAATDAT D	AATDTATĐAT	AATOTATOAĐ	TTATAAAAA	DAADDTATAA	746621
ADTAAAAATA	AASTSSSTTS	ATTATADATA	GCAAAAATGA	TDADAADDTT	DTTTTAATDA	
DTADTTTAAD	TDATACATAA	ADSTDTTTTA	DDDAAADTTA	TAADABATBA	AATADTADDA	105911
AAAAAAAAA	AAADAAADAA	DDAAAAATDT	ATOTOODADA	DYDATADAAD	CCAGCCTGGG	T##9#T
ADADDTTDAA	TTGAGCCCAG	TOTTADDADD	GGCCAAGGAG	AASSTITSSS	AADDTATADA	146381
				AADDTTAADD		176351
				AGTAADAADA		19291
				DAAADTAAAA		146201
AAADAĐĐTAT	ADATATAADT	DTDTDDDADA	CAACAGAGCA	DDDTDTDADD	TOADSTOADD	175971
DTTDTADTDD	TGCAGTGAGC	DDDAADTDAD	DDATCODADT	TOADTADDAD	DDTDDADTOD	180971
DADDITITAD	DDADDDTTAT	DICCETGCCTG	DOTODIDIO	ADDADTAAA	AAAADATAAD	146021
TOTOTOOOAA	ADADDTADAA	Seestage	SASAASTTSA	TOACCOACT	TODITIADDAD	T965#T
DDTDDAATDD	DADDDTTTDA	DAADDDTAAT	DTDDDAADTD	DOTODIACOD	ADDDAADATO	T065#T
					1000440440	100311

152381	TGGTTGGAAG	AGTCAATGTT	ATTTTGATTT	TTCTGTTTTG	Thurst Carrent w	A A MOCO A COMMO
152441	GCGGYIWYII	CCAGCITICI	TTCATTCCCT	. ACATGAGTTC	AAATCCCACC	3336555
152501	GONGANCUCA	GACCTTCTGA	L CTTGTGGGT	. CCCCTDCTC	TORCOTORS	
152561	***	. IGACCCATTA	AAGACGGATC	GAGACAGCAA	CATACCAMON	mas
152621	C110C1110C	CCCAGICCAG	GTTAACCATC	TGTGGTATT		10000
152681	CONCATAM	NICAMITATA	TATCCACTAA	AATCTCAGCA	CTACTCTARG	M1 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4
152741	ATGACAGCGA	AGAAAACAGA	CCAAACGTCT	GCCCTTATGG	CIAGICIAAC	TACTAAGGAA
152801	TGCTGGTTAA	ACCAAGGAGC	TTCTGCTCTT	TTCCTTAGTC	ACCTCCCCC	ATTTTCTCTG
152861	AAGGAGAATA	TTGATAAACC	TGGAAATAGG	GCCGGAGAGT	ACCIGGGGA	GGCAGAAACA
152921	GGAAAGTAAA	GATGTGGCAG	CCAGTATTCC	CGTTATAAAA	GGATAGAAG	GAAGCCTTCG
152981	AGTCCAGAAA	AATTCCCACA	AGCAGGGGCT	GCTCATGCAG	ATCARCARCT	CCGGCCTCAT
153041	AAGTAAGTGC	TACATAGCCT	TTCTTTTCC	ACAGCCTGAG	CCTCCACAA	GTTGGGGGAG
153101	GCTCTTGCTT	CATGCCAGTG	CCCCTCTGCA	CATTTTCCAT	GGICCAGAAT	CCAGACTGAG
153161	CGGTTCCTTC	GCCAACATCC	ACTTCAAAGT	AACGTCTTCC	MCAAACTCCT	AAATCCCATC
153221	CCAAGACACA	GGGGAAGGCA	GTAAATCTCC	TGGAAGATGT	CTCCTCATT	CCTTCACAAC
153281	ATCCACGAGT	CACTTGTCTC	CGATCCTCAG	AGAGAATTAG	GICCIGATTC	TCCTGGGTGT
153341	GATCCAGAGT	CACACTAACT	GCDADACADA	ACAAAACAAA	TICGIGATGA	GCTGTATCTG
153401	TGAAGAACAC	AGGTTATTTT	A THEFT ATTEMA	ATTTTGAGAT	CAAAAATAAT	TTTGTTGCTG
153461	GCTGGAGTGC	ACTGGCACTA	TCTCAACTCA	CTGCAACCTC	GGAGTGTTGC	TGTCACCCAG
153521	TTCTCCTGCC	TCAGCCTCCG	GACTAACTCA	GACTACAGGT	CACCTCCTGG	ATTCAGGCAA
153581	AATTTTTTTA	AATTTTCTCT	AGAGATOCCC	TTTCGCCATG	GCGCACCACC	ACAAGTGGCT
153641	CTCCTGACCT	GAAGTGTTCC	ACACATGGGG	TTTCGCCATG	TTGGCCAGGC	TGGTCTCAAA
153701	GAGCCACCAT	GCCCAGCCAC	ACCUACCICG	GCCTCCCAAA CAATAAAACC	GTGCTGGATT	ACACAGGTGT
153761	TATTGTTTCT	TATAAACTGG	GTGACCTTAC	GCAAATCATT	AGCCTGTGTT	CAAACCCAAC
153821	TGTTAACTAT	AAAGTGGAAA	TTACCCTATT	TGTTGCAGAG	TAACTTTCTG	AGCCTCAGTT
153881	AAGCTTATGT	TTGCTTAATG	CTTGGTNAN	TTCCTGGTAC	AATGGTGGGT	AGGATTGAAT
153941	GGTAGTTGTT	GGGGTGATCA	GCCCAACAC	CAGGCCGTGG	ATGGTAACCA	CCTAATAAGT
154001	GTCAAAGGAA	TGAGAAAAGA	CARCTERACAC	GTGCATAAAG	GGGCTACAAA	GTCCGGCGGG
154061	TAGATTGGAG	GCTGCAAAGG	CCCTAACCTC	TGGGAGCCCA	TGGGTCCAGG	GTGCCAGCAC
154121	ACAAAGAAGC	AGGTGGTGAG	CACCTCACCC	TOGGAGCCCA	CACTATTTAT	TGGTGATCAA
154181	TAGAAAGGTA	GTGGTGCATT	AACCCTACCT	TAAACAGGTG	AGGGCATGAG	GACATGGGGG
154241	TAGAATATAC	TOTGOTGOTT	GAGATACTAC	GTGACAGTTT AGGACACGTT	AGCATTTCT	TTGACACATG
154301	CCAACAAGTC	TGTGCACTTT	CCACACCCCA	TGAGGGGTTT	TATGAGTGAA	AAGCAAGGAA
154361	CCATCCAAGC	CACAAGGGGT	TTTNTCCCCT	AGGCTTAGAT	TATGCCCTGA	GCCCTGGGTT
154421	CTTCCACCAT	TTGGCACAGA	CCTTCCTCTT	CCAAAGGCCA	TTGTGGTGCG	GCAGGGCAGC
154481	ACCCCGGACA	TCTTCCAAGA	CTCTTTTACA	TTATCACACA	CGAGGGGTTT	TGGACCCTGG
154541	TCTTCTAACA	ACATGTAGTA	ATANTONTAT	TTATGACAGA	CAAGCCAGTC	CTGCTTCAGC
154601	AGGATGCCAA	GGTACAGAAC	TAACCTCTTA	CATCAACATC	ATCTTCGTCT	TAATTATTCA
154661	TCCCATGCAG	GACTTCCAGG	AATCATCACA	ATATGGTTAC CAGTTGAGCA	CATCCTGTCC	AAAGTTCTTC
154721	TCTACTGAAT	AACCACCAAC	ATTCACAAAGA	AGAGAGGGAA	JAAAGATACC	TTTTCCCTTC
154781	AGCTTGTTAT	TGGAAGACCC	ACCTCTCATC	ACACATGCCT	AATGACTCAG	CTAATGTCTT
154841	TAAGCTCTTC	TCTTTCCCCT	Chantarer	TCCATAAGCA	AGTCCCATGA	CTTTTAATTG
154901	ACTGAGGACC	AATATACATG	ADDDDTATCD	GACTAGAATC	L'IAGTATGAG	ATAATAATAC
154961	CTGATAACCT	AAAGTGAGAT	ACTCAACACT	ATGCAGTTTT I	AAACAAGACA	GAAAAAAGAT
155021	GGATGTTCTA	ACAAGAGAGT	TARCARACAGI	CTGTGCTACT (AAAAATAAAA .	AATGGTAATA
155081	GGTCTGTGAC	AATTAAGGAA	TTCAACAACCA	CIGIGCTACT (GAGTTAAATG	ITGATCAGTT
155141	GTTTGTTCTT	ררב מדממדר	CCTCACGIAII	CAGAAACACT	CCTGTGCTG (GATGCTCTCT
155201	ACATGGACAG	ATTABCCACC	CTTTCCCCCC	CCCTGTCTTG (AGGAAGGCTG
155261	AGAGGAGACC	ATAGGGGAGA	PACANCONCO	TCTGGCTTGG 1	TCAGCCAAT (GGGAAGCACC
155321	CTATGATTTC	GAGGGTCTGC	ATTCCTCTCC	CTTGGGAGTA 1	TCAGTACCC (LAGTCCCACG
155381	CCCTACACCT	GCCACTTCAC	GCCCAGACCA	CTCTGGGCAC A	ACTOTAGTAT A	AGTTACAGCT
155441	GGGTGCTTCC	TGTTCCTTGT	GCCCAGAGGA	ACTOCTOR OF	TCTAACTGT '	CCTAGTTCT
155501	TCAAACTCTA	TTCAGTTAGC	GGAIIICCCA . TTTTNTCNCC	CTCACTCACC 7	TTGTAAATA (CCTCCTTT
155561	ATATTACCTG	AATGACCCAC	TITIATCAGC	CIGACICACA (MAGTITGGG (FITTCAATTC
			GAMMACCCAT	GIIGAGAAAT 1	AAAATGTTT)	ACGGGGTGGT

Figure 9 (Page 48 of 74)

155621	AATACCACTT	AAGAGAAAA	8 T 8 T C 8 8 T T C	Characannan	ATTCCACCTA	777 P
155681	TGACACATCA	ACAAAAACAT	ATACAATIG	TCCAACCTAA	AAGATAGATA	TCTATTGGTG
155741	ATACTGTTAT	ACTATTATAT	CARAGAMAGAI	TARCTCACAC	CATTATTA	AATGGAAGAA
155801	GGGCCAGGTG	TEGTEGETEN	TOCOTOTANT	CCCACCACT	TGGGAGGCCA	AATGGAAGAA
155861	ATCACTTCAA	CCCACCACTT	CARCAGO	CCCAGCACTI	TGGCAAAACC	AGGCAGGCGG
155921	CAAAAATACA	ACAATTACCT	CAAGACCAGC	CIGCCCAACA	TGGCAAAACC	CTGGCTCTAC
155981	CCCTCAACCA	CANCANTAGET	GGGCATIGIG	GCACATGCCT	GTAATCCCAG	CTACTTGGGA
156041	CCCACTAGLA	CAAGAATCAC	TTGAACCGGG	GAGGCAGAGG	TTGCAGTGAG	CTGAGATTTC
156101	GCCACTACAC	TACAGCCTGG	GTGACAGAGA	GAGATTCTGT	CTCAAAAAAA	AAAAAAAAGA
	AAGAATGAAA	GGAGTCACCT	AAAAAAGATA	ACACAATTTT	AAACATAAAT	GTACTACATT
156161	ATTAGTGAAT	TCATGTTTAG	AATTGTGTTA	ATATACAAAG	CAAAAATTGT	AGAATTATAG
156221	GAGAAATGGA	CAAATCTACA	ATCATCATGG	GATGTTTTAA	CATTCTTCTT	TCCATAATTG
156281	ATAGATCAGG	CAGACCAAAA	GAAAGAAATA	AGGGAAGATA	CGGAAGGTCT	GAACAATCTA
156341	AGAAGCGCAA	TCTCATAGTC	AATACATAAA	GCTCAGCAAT	TGTTTAATAA	TAGTAAGCAG
156401	AGAATATGCA	GTTTTCTCAG	GTATAGATGG	AACATGCACT	AACTGAGTAA	ATACTAGGCA
156461	GAAAACAGTC	TGAACAAGTT	TCAATAAATC	TGTATTACAC	AGATCATTTT	CTCTAGCCTC
156521					AGATTCTAAA	
156581	TGTAAACTAC	TAATAAGTCA	TTAGAAGATG	TATAGAATGG	AACAATAATA	AAATGTTATT
156641	TATAAAAATA	TACAATGAAG	CTAAAGCAGA	ATTTTAAGGA	AAATTTGTAG	GCTTTAAATG
156701	CTTATCTTAG	AAAAATTAAA	AAGCTGAACA	TTAATGAGCC	AAGCATCTAA	TTTAAATTTT
156761	AAAAAGAACA	TAGAAAGCCA	AATATAATTT	TTTAAAAAGA	AAAAATAGAT	ATTAAACAAT
156821					GAAAAACAAA	
156881	AGTAGCTTCT	TTTAAAAGAA	AAATAATTAA	ATAGACATAC	CTCCAATGAG	ATTTATCAAA
156941					TTAAATATTA	
157001					AAAATTATTA	
157061					GCATGGGCAG	
157121					ATAAAACGTG	
157181					AGTCAAGGCA	
157241					CTCCCGCTTA	
157301					ATTTTATATT	
157361					TATATAATAT	
157421					ATATATAAAT	
157481					TAATACATTA	
157541					AATCCCAGCA	
157601					AGCCTGGTCA	
157661					TGTGGTGCAC	
157721					CTGGGAGGTG	
157781					AAGGGAGACC	
157841					GTAGTCCCAG	
157901					CATGATCGCA	
157961					ACAAATATAT	
158021	ATCAATATAT	ATATTATATC	TACCANTCAN	TECTTENETT	TTATATATAA	TATACACACIA
158081					AGATACAGAT	
158141					TATAGCATAT	
158201					TGCCTGTAGT	
158261					TGGTTGAGCC	
158321					GTGGAAGGAT	
158381					GTGGAAGGAT	
158441					TGTGTGTATA	
158501					GAGTCCAGGT	
158561					CATAGTTACG	
158621	GTCTCCAATG					
158681					TCATCATCAT	
158741						
158801					TCTTCTTTCT	
73000T	CTCCCCCACC	ICITICTUTT	CCTCCTCCTC	CITCATCTCT	CITCITTTT	TITTTGAGAT

Figure 9 (Page 49 of 74)

158861	GGAGTCTTAC	TCTGTCGCTC	AAGCTGGAGT	GCAGTGGCAC	AATCTCAGCT	CACTCCAACC
158921	1010001101	I GGGTTCAAGC	AATTCTGCCT	r Aageeteeag	AGTAGCTAGG	NCTCCB BOTT
158981	CACACCACCA	A CACCIGGCIA	ATTTTTGTAT	מסמדטמדדדד י	CATACCCTTT	C. C
159041	GCCAGGCTGC	- TCTCAAACTC	CTGCCCTCAA	GTGATCCTCC	TGCCTCGGCC	TCCCSSS
159101	CIGGGATIAC	. AGGCGTAAGC	CACTGTACCC	GGCCTCCTCC	TTTAATAGAC	ACCCCCCC
159161	ICIGIIGCCC	AGGCTGGGTA	CAGTGGCGTG	ATCATAGCTT	ACTGCAGCCT	CCARCOCC
159221	GGCTCAGGAG	ATCCTCCTGC	CCTAGTCTCC	CCAGTAGCTG	GAACTACACC	CGAACTCCTG
159281	GGGGCTAATA	AAATTAATTA	GGTGATAAAA	TTCACTGCCC	ACTGATGACT	ANAGCACAC
159341	GACATAAAAG	ACACAGACCT	TGAAGGAAAA	TGTGTCTACT	TAATTTCAL	AAGCTCTTTG
159401	TCAAAAAAACA	GGATGAAAAT	GCAAAATGCC	ATCCACATGC	CACAACATAT	CLCCTATTTA
159461	AAGTTCCCAT	AAATCAATAA	GGAAAAGAAC	CCAATAAAA	TTRTTRACA	CAGCTATAAT
159521	ATGGGTAAAT	CACAGAGGCC	TGAAGGGCTA	ATGGACATAC	AAAAAACC	ACAGTAAATC
159581	TAGTGAAATC	AGAAAAGCAC	AAATTAAGTA	CACAATTAGG	TACCATTTTA	TCAATCTCAC
159641	ACTGTCAAAA	TCATAAATTA	TATAAGTAAA	GACTCAGGGA	GTTTTCCACC	AATCTGTAAG
159701	CTTATATTGC	TTGTGGGGTA	GAATTGGAAC	AATTTCAAGA	TCTCTACTAT	AGIGAGAGCT
159761	TATGATATGC	ATCCCTCACA	CCAGCATGTC	ACTCCAAGGT	ATCTCCCTCC	CIGGIAAAAT
159821	TACGGGACAC	AAGGAAGCAT	GGATAAGAAT	GTTCACAGTA	GTATTCTCTC	AGGGAACATT
159881	AACAACAAAA	AAACCCAACT	ACACACAACT	TCAATGCCCA	GTCCACAACC	CAACAGCAAC
159941	AATAAACTTC	AGGCCGGAGA	TGGTGGTTCA	TGCCTGTAAT	CCCAACAAGG	CAATGGATTA
160001	AGGCGAGAGG	ACTGCTTGAG	CCCAGGAGTT	CAAGACCAGC	CTCAACACII	TAGAAGGCCG
160061	GTGTTTCTAC	AAAAAATTTT	TAAAAAATTA	GCCAGACGTG	CLOWLCWWW	TAAAGAGATA
160121	CAGCTACTGG	GGAAGCTGAC	GTGGGAGGAT	TGCTTAAGCC	CACCAATTO	CCTGTGGTCC
160181	GAGCCATGAT	GGGGCCATTG	CACTCCAGCC	TEGETERANCE	CAGGAATTTA	AGGCTGCAGG
160241	AGATAAGTAA	ATAACAACTT	TGCATTTTCT	GCCACATTCC	AGIGAGACCC	TGTCTAAAAG
160301	TCTAGACTCT	AGACTCTTTC	TATGACTACC	TTCTACTTAT	CACATCCTAC	GAGAGTGGTT
160361	TAACCTCTCT	GTGTCATATT	TCCTCCTCTA	TABACCAAAA	ATCCCCCATA	AACACTCACC
160421	TGTGATATAA	AACAAGAACC	AAGAAAAGTA	AAGCTTTTCT	AIGCCCCAIA	TAGAGAGGAC
160481	GTGCTCAGTA	TATGTGAGTC	ATTATTCCTG	GTGCTGGTAG	CACTCTATCT	AGACTAAAGA
160541	AGTCAAGTAA	TATGGTACCA	TATATTAAGA	TTAACAACAA	CCTCCCCNAT	COCACCTTG
160601	GGTATGTTCC	CAAAAGAAAT	GAAAGCACCA	GGATATAAGG	ATCCATCCAC	CCCAGITIGG
160661	TTGTAGCAAC	ATTGTAATAA	CTAAGTTCTA	DOMINIAGE	GARCTTCCAT	CAGEAGGE
160721	ATGGTTACAT	ATATTTATTA	TATTCTTATG	GAATATTAGA	CATAAAAAACT	CAGTAGGGAT
160781	ATAGAAGAGA	CAGTGTATAT	ATGTTACGTT	TGTACAAACT	TAGGGAAAGA	TATACATTAC
160841	CCTACCTAGA	GAAGTCAGAT	TGGAGAGGGG	TGGGAAAAC	CTTCAACTTT	CTCCTTA
160901	CCTTTATATT	GTTTGACTGA	TTAAAATGTA	TTTGTTGCAT	CTGCTTCAAC	CCCTTATAT
160961	ATAAAATAAA	CATACATTTA	AAAATAAAA	TAAAATTTAT	TCCTATCACT	GCAATGTAAA
161021	AGCTGGGCAC	AGTGACTAAC	ACTTGTAATC	CTAGCACTTT	GGGAGGGACA	CACACCACA
161081	TCACCTGAGG	TCAGGGGTTT	GAGACCAGCC	TGGCCAACAT	TOTON NOCCO	GACAGGCAGA
161141	AAAAATACAA	AAATCAGCCA	GGCATAGTGG	TGCGTACCTG	TARTCCCACC	CATCTCTACT
161201	GGCTGAGGCG	CTGGAACCCA	GGAGGCAGAG	GCTGCAGTGA	CCTCACATTC	CIACCCGGGA
161261	AGCCAGCCTG	GGTAACAGCG	AGACTCCATC	TCDADADAA	ATTTCAAAAA	ACA A A A A A A A A A A A A A A A A A A
161321	TAATAAACAG	TGTTTAAGAG	GGGAGAAATA	TTTAGTTAAA	AGATAACCCC	AGAAAAATTT
161381	TAGTTTCACT	TGACCCGGAA	GGCGGAGCTT	GCAGTGAGCC	GAGATCCCAC	CACTCCAACTC
161441	CAGCCTGGGC	GACAGAGCGA	GACTCTGTCT		ANDRICGCAC	CACIGCACIC
161501	AAGAAATAGT	TTCACTTGAA	CCATATTATG	ATTCCTTCTC	TABBACATCA	CACAAGAAAGA
161561	ATTGACTCAG	TGAAATCCCA	GCAAAACTTA	CACAAAGTCT	TGTTCTTCCT	TCCTCTCARC
161621	TGTATAGGAT	GAAATACAGA	GTGCTTTTGG	GTTTTGTTGT	TGTTTCTTCT	TCTCTTCATC
161681	AGGGGAACAC	AGGTCTATAA	TTCCTTTTCT	GARATCCCTG	CAACAAAATC	CCCTTTCCC
161741	TTCAAATTAG	TTTAGAAGTT	ATAAAGGCAA	AAAAATGCAT	ATACTCTANA	GGCIIIGCCA
161801	ATCATGGCCT	AAGGCAGAGC	CCTGTAATCA	AATTCATCAA	TATATOTOTOON	GLICAACCCC
161861	TATTCAAATT	AAGTGGGATA	AATAAAGACT	TTTAAATAGT	CTCATCTGCA	TGCCGTTCAC
161921	GGTTGGCCAC	TGTGGAAGAC	AGACTCAAGG	GTGGCCTTCT	ATGATTCCTC	CCTCTTCAG
161981	TTCACACCCT	CGTAAAATTC	CTTGTCTTTC	AGTGTGAGCA	GGGCTTbTGA	ATTGCTTCTC
162041	ACCAATAGGA	TATGGCAAAG	ATGATGGGAT	ATAATTTCTA	TGATTACCTT '	TCDTTDTCTA
					m MCG11	LCALIAIGIA

Figure 9 (Page 50 of 74)

162101	AGACTCCATC	TTGCTGGCAG	ATTTTCTCTA	AAGAGTCTGT	CTCCTGAGCT	CTCTCTGAAG
162161	AAATAACTGG	CCATGTTAGA	AGCCCATGTG	CAAAGAGCTG	AGGGGTGGCC	TGTAGAAGCT
162221	GTGGGCAACC	TCCAGCCAAC	AGCCAGAAAŤ	AACCAGGGCC	AAAGTCCTGC	AACCATCAGG
162281	AAAGAAATTC	TGCCTGCTAT	CTCAGTGAGC	TTGGAAGTGG	ATTCTTCCTT	AGCCTAGCCT
162341	CCAGATAAGA	ACACAGCCTG	ACCAACACCT	TAACTGCAGC	CTTATCAGAC	CCTAAGCAGC
162401	AGGCCCAACT	AAGCTGTGCC	CAGATTCCTG	AACCACAAAA	ATTGAGATAA	CATATCAGTG
162361	TTGTATTAAG	GTTCTAAATT	ATGGTAATTT	GTTTGTACTA	ATAGATAACT	AATATAACCA
162421	CCAAATCATT	TCAGGTTAGG	CCAGATTTTT	GTAGCCAAAT	GAATCATGAT	AAAACTTTCC
162481	ATTTTCAGGG	GTTTTTTTGA	TTTTGTACTT	ACGGATACAA	ATTTGTGAAA	GTATAGTCAG
162541	CACTGATTTA	AAAAATCAAG	GGAGCAGGAA	ACTCAGTAAA	TGGTTCTAAC	ATTTTGGAAT
162601	CTGTAAATTG	GTTGTAACAT	TTGTCATCTG	TGTTATCTAA	GTCAAGTTCC	TAAAATATGT
162661	GAATGATAGG	TTATCATACT	CACCTACTTT	TCTTGCATTG	CTCTAAGAGT	TGGCTGAGCT
162721	ATTGATAATA	AACACTATGA	TCAGATCTAA	TACCATGATG	TGCTATTATG	ATCATGTGTC
162781	AGTCACAGGG	CTAAGCACTT	TGTACATGTT	GATGCATTTA	ATTTTGATGA	TAACTCAATC
162841	AAGTAGGAGC	TGTTAATATT	TTCATTTTTC	AGAGGGGGAA	ACCAAGTCAC	TTGGAGTAAC
162901	ATGGCTAATA	AGTGAAAGAA	TAAGAATTTG	AAAGGTTTGC	ACAGATAACC	AGAATGCAAT
162961	GCTCATCACA	TTCACTGAGC	AGTGAATCAT	ACTAACTAGA	GAAAGTATGA	AACCTCTACT
163021	GAAATTAACT	AAACAACCTC	TCTGGCTGTG	AGCCTGCCAA	GGGACAGGTG	GTAAACTTCC
163081	TTACTGCATA	AGGCCCCTTC	TATCCACAGT	ATTCAGGAAT	TCTTTAGTGA	ACATACCTTC
163141	ATGACTCCTT	AACATTTTCT	TCACATCGAA	GTAAAGCTTG	GAAACATTCC	ACATACCITG
163201	AAGTTCCAAG	GAGACAGCCT	CTGATGTTTC	CAGCTTCACA	GCCCAACTCC	TACATAGIAIG
163261	AGAGGCGAGA	GATTTCTTCA	GAGGTGCATT	CCATTCATTT	CTATATACCC	ACACCCCTCC
163321	CCTCCTGCAT	TCAAACAGGA	CTTACCTGCT	CAAAGTGTCA	TTCACATTCT	ATABACAAAC
163381	AAAAAGAAAA	GGTGAGCATG	GGAACATCGG	TATTTCATGG	GGCTTGTCAT	GCACCCCTAT
163441	TCTTCTTTGC	TTTACCCGAA	GAAGTAAAGA	GAGTTACCCT	AGTCTTAGTC	TTACATATTC
163501	ATGGATACTC	AAACAAAGTA	ATTCCCACCA	GTCTTAGGTA	TTGATGGATA	CCCACATCCA
163561	ATAATTCCTA	CCAGCTTCTG	GGAGATTCAG	CATGGCAGGA	TGTTTATCAA	COCAGAIGGA
163621	TATTCTCATC	CTTGCTGAAG	TCTGAGGGCC	AGGAGCTTTG	TCCATGCTCC	CTCTCTAACC
163681	ACTAGCTTTT	GGTGATCGGA	TTTCCTTCAC	AGTGAGCCCA	GATTAGAGAA	CICIGIAAGG
163741	AAAGGTCCTT	AGTGGTGAAT	CTGTGCACAG	CCCTGAGACT	GGGCCACTGC	CACTIAICAI
163801	GTGGTAGCAG	GTATCACACA	GTGGTAAAGC	AATCATGCTA	TACACTCAGC	CTTDCACTAT
163861	AGTCACCAAT	CCTGTTAGTT	AGAACCAGAA	TTAATGGCTC	CAGATGTTTA	TCTTCCTACA
163921	GATAAAGCTG	TAGATTGTAC	CATAACAGCT	CTGGAGCAAG	GGTTCTACAA	CCAAATCACA
163981	GAAAAGGTTA	TCACTCATTT	TGGCTGCCCC	ACTTCATCAC	CCATCACTCA	CCTACTCCAC
164041	TATTTCAGGA	GAGAGTCAAC	AACCAGGGTT	CTCTGCACAT	GGGCCAAGCA	CCCAAACACT
164101	GGTAAATGTT	ATCCCGTGGT	TTCATTTGGC	CAAGCTGTGT	TCCCTCACAA	CTTTA TTTTT
164161	CTAATTGACA	TAAAGGTACC	CTATAAATTA	GTGAAGGCCA	CCCTCAGAA	CTCATCTT
164221	ATCTAAAAGA	AACATTACTT	TATCTTCCCA	TECTTCCTTA	CCATTCTCT	TTAATAGGAG
164281	TATAACATAC	CTTTTTTCCC	TACTCCAAGT	ACACAGCCTC	ACCTGCAGCA	ATTTCTCCCC
164341	TGAGCCCTGA	CATTTTTCCT	CCAGTTCCAG	GATGTGGCTC	TTCACTTCAT	TCCTCTTCAC
164401	CCCCAGACCA	GCCTCATAGT	CCCTCAGTCT	ACTCAGAGTC	TCTTCTTCTT	CTTTCTCCAG
164461	CCTCCAGAGA	TAAGACTTCT	CTTCCTCATG	TAGGAAACAC	TGGAGATTCT	TARRETCAG
164521	CCGGATTTTT	TGTCTCTGAA	TCTGTACCTT	CTCCTGGAGT	CAAGAAACTA	TCCTCNANG
164581	GTGGAAGTAA	ACCAAATGTC	CATCTATGGA	TGAATGGATA	AACAACAACIA	AAACTCTCAC
164641	ACACGCTACT	ACATGACAAG	CCTTGAAGAC	ATTCAAGCAA	AATAAGCCAG	AAAGICIGAC
164701	GCAAATATTG	TAAGACTTTG	CTTATACAAG	GCATCTGGAG	TACTTAACTT	CATACACACA
164761	GAAAGTAAAA	TAGTGGTTAC	AAGGTGTTGG	CAAGACCAGA	AATTCCACAC	TTATTCTTTA
164821	ATGGGTAGTG	AGTTTCAGTT	TAGAAGATGA	AAGATGAAAC	TGAGTTGGAG	TTTCCACATC
164881	GGAATGGTGA	TGGTTGCACA	ACAATGTAAC	AATGTAAAAG	CACTTAATTC	TACTERACTA
164941	TATACTTAAA	AGTGGTTAAA	TGCTTAAGTG	ТТАТАТАТАТ	TTTCDCACAA	TACIGAACIA
165001	ACACACAATC	AGCCACTGGG	ACATTATTTT	CTCATGAGTC	ACTGARGCTC	CA ACA ATCTC
165061	CCCAGTTTCC	TGCTGCAGAG	TCATGTGTGG	GAGGCAGGCA	CTCAGATGTG	GAAGAAIGIC
165121	CCTCAGATTC	CTTATAGTCA	CCCAATTAAT	TTTCTTCTTC	TTCAGCCAAC	ACACACCACA
165181	AAGCTGGGTT	AGGAGTGCTA	GATAATTTAA	TTGTGAAACT	AGGGCCAAG	TCA A A CA CTT
					LUMMATABLE	ICAAACACII

Figure 9 (Page 51 of 74)

165241	TATCAGTTAC	AAGGATAAAA	AGAGGTTTTT	ACTTATGATT	TAAGAAGTTA	CATTTCTCA
165301	LIGGAGCGAT	TITCTTGAAG	TAAAAGCTTA	TAATGAACAT	CACCCACACT	CCAMMMAAA
165361	ACAACCAGGC	TGGTAAGAGG	GTCCATAATT	CTTGGCAGGG	GGAGCTTTCA	CTCTCLCLC
165421	CATTIATIAL	GGITAACTGA	GAAATACTGT	TCTACTACCC	TAGGGTCATC	TTARCCRITTO
165481	CIAIGIGIAA	I GACTGACAGA	AATCAAGTGA	AACTCTCATC	TGAGGAGATG	TRARCTTCCA
165541	ATTICCATTA	GIGCIGICTA	AATTAATGCA	GTGGGAGTGT	GTATTCAGGG	CAATTTCAAM
165601	CTATGTTCTT	GGATTGCAGT	CTTCAAACTT	GGCCCAAATA	AACTCTCTAC	TENTOTERA
165661	AAAATAAAA	ATAAAAAATA	AAAATAAATT	CATACAGTGT	TTTGATGACT	ATCATANA
165721	AGAAGGGTCT	TTGACTTAGG	ATGAGGTGGA	ATTTTTGTGT	AGGAGACAGG	TECACETTE
165781	ACTCTTGTAT	AGACGGGTTT	TCATATATGT	TAGTTACAAT	CAAGGTCTTC	CCCATTCCCC
165841	AAGATCCTAG	AAATGGGGGA	AGTAAGAGTG	TACTCAGGAG	CTCAAGAGCA	ACATIOCOC.
165901	ACAAAGATCA	GGGTAGAGGT	TAGAGAGGAC	TCCTGAAAGA	GAGAAAATTG	GTAATCACAA
165961	TGTGGGATTT	TACTGCAAGC	TAGTGAATTA	TATAAATATA	AAGATTGGTG	CAAAACTAAT
166021	TGTGGTTTTT	GCCTTTACTT	TAATGGCAAA	GACCGCAATT	ACTITICAC	AAACCTRAAM
166081	ATTTCCATAA	AAGAATGTGG	CTCTGATAAT	GTGGAGGTTA	GTCNGCCNCC	CAAACCIAAAT
166141	GAAAGTTTGT	AGTTGCAAGT	GTGTAGGTTG	TTGCATTACT	TGTCATCTAG	GAAATAATCT
166201	AGTATAGGCC	GGGTGCAGTG	GCTCACGCCT	GTAATCCCAC	CACTTTTCCCA	TIATAAATCA
166261	GGTGAATCAC	GAGGTCAGGA	GATCAAGACC	ATCCTCCCA	ACATOCTON	GGCTGAGGTG
166321	TACTAAAATA	CAAAAAATTA	GCCAGGCATG	GTAGCACATC	CCTCTAATCC	ACCCCGTCTC
166381	AGAGGCTGAG	GCAGGGGAAT	TGCTTGAACC	CGGGACCTCC	ACATTECO	CAGCTACTCA
166441	CGCACCACTA	CACTCCAGCA	AGACTCCATC	TCDAAAAATA	ACATIGCAGT	GAGCTGAGAT
166501	AAATAAATAA	AGTATATTTC	TTTCATCAGC	TTCATCACCT	GIAATAATTT	AAAAATAAAT
166561	CTGGAGTGAT	CCTGTTTTCT	AAGTGTTCAC	A A CCCTCCT	TGAGTAGTAT	GAATTTCAAT
166621	GAGCCAGATG	CTCCACTGTG	GTAAAAGTGC	CACCCTAATC	POTTCACCT	GTAAAGTTGA
166681	GTTTATTTTG	AGGTATTTAA	ACTTTCACAC	CAGGGTAATG	AGTTGAGGCC	TGCAAACCAG
166741	ATACTAATTC	TGCTTCTTCT	CACTCAACTA	TCACTCGATG	CTTTTTCTAG	GTAAATAGTC
166801	ATGGAAAGAT	TGGTGCTAAA	TACTCATCCA	TCAGGAATCC	CAGCCAACTA	CAGTTTAAAG
166861	AATAATGGTT	TCTTCCTTGG	CTUTCATOOA	TTCAAACCIG	GAACCAGGGG	CATAAGTACA
166921	ATTGTGCTTT	TCCTCAATCA	TCCCCTATGC	CTAACCTCTA	CANTOCARA	ATAAATCCTC
166981	TCAATGAAGT	CAGATTCTTA	CTTTCCATTT	ACTUATOR	ATTCCTCTC	TAGCTTGAGA
167041	CTCCGTACAT	CTGTCTTCAA	GTTGCTTCAG	TTTTCTCACA	ATTGCTGTGG	ACAGCTTCTG
167101	AAGGAAAAAT	TTGATAAGTG	AAGCCTATTC	AATTTCACTC	TTCATTAGA	COTTTCCTG
167161	AATCCCAATC	TTCTAAGATA	TATTTCAATA	ATTICACIO	TTTTATIAGGG	ACCTAGGGGG
167221	TTTTGCTAGA	GAGCATGCTA	AAGGCTATAT	CTCCACCAAA	ITTATAGAGT	CCTCATTGTT
167281	CCTGAATAGT	TGGTAGGATT	TTAAACTTCA	TOTAGGAAC	ATACTGATCC	CCTTGGCAAC
167341	AGGGGTAAAA	TGGTAGGATT TAACTTGCCC	AAACTICA	CACTOCCT	GTAGAAAATG	AGACTAAGAA
167401	TCTCATCTCC	TGACCCAGAG	CCTCACCTAT	GACTGCCAGG	TGGTGGAGCA	ACAATTGCAA
167461	TTGGATATAG	AACAAGGTAA	TCATCATCTA	DARACTE	AGAGTCCTGC	CAGGAAAAAG
167521	AGCAAAACCA	ATACCAGTGT	TTGGCACACA	TCAAATTTTC	TAAAACAACA	TGCTGAACCA
167581	ATCAGGATGC	CAGCTGGTTA	TTAGAAACAG	TTCATCCAAC	ACCCCAACCC	GTCAGGAAAA
167641	TGAACAATGG	TATCATGAAT	CCDDTTTDDD	ATC A TOGANG	AGGGGAATTC	TGGTATCTTT
167701	TTATTCTTCA	AAACAGTTTC	TCATATTTCT	ATTCARROT	ATTCATGTCA	AGCTTTTAGC
167761	GCTAATTGTA	GTCAATGCTG	DADGATTCT	CTCCTCTCCT	ATTTGAAGCT	GACCCAAATT
167821	CTCATTCATT	CTCGAGTGTT	CTCACCAAAA	CTCCTGTCCT	CIGIAAACCC .	AACAAGTATA
167881	ACATTGTCCT	TACTATATGC	CICAGGAAAA	GGTTCTATGT	AACTGTTTTA	GCAAAAGATG
167941	AGCTTATAAC	CACCTCCTGT	CTATCTCTTT	TREGGREGATI	CTATATTTTA .	ATGTCCTCAA
168001	TTTACAGATG	CACCTCCTGT GAGAAACCAA	GCTCTCAACA	AGGGAGGGA	GGACACTGCT .	ATTATCCCCA
168061	TAGTAAGTGA	CDDDDCTCNA	TTTCXXCX	ACCTOCTOC	CGTGCCCAAA .	ATTGCCCATC
168121	AAAGTAATTC	CAAAACTCAA AAATGGGAAT	ATCAMCATA	AGC I GGTTCC	TITICTTACT .	ACTTGGTGGA
168181	AAGAGCTGCC	ATGAGCTGAG	TEGTECTONT	CAGTTATTAG	CIGCTCCATG (GAGTTTAAGG
168241	CTTCATACAA	GACCACCTOAG	GCCTCNTCCT	CCACACATG	ACCATAGAAG	GACTTAGAGC
168301	AACATTTTCC	GACCACCTCT TCAAATTTAG	GCTCATGGA	GGACAGAATA .	AGGAGCCTGA	CACTGGAGAC
168361	TTCCTCCATC	CTCCCAACAC	CARGGACAGA	COTTCOTT	GGACATCAGG	ACTATGCCCA
168421	CTGGATGGTA	CTGCCAACAG	CTCCCTCTCC	CCTTCCTTAA '	TATGCTTTCT (GGCAAGAAAT
	AIDDIAGIA	CACAAAACCT	CICCICICIC	LICACCTTCC .	ACAACCAAGC /	ATTTCCAAAT

Figure 9 (Page 52 of 74)

168481	CTTTGACTCT	TCTTCCTGAA	TCGTGCTTAA	AATCTGCCCT	CTCCTCCCTT	TCTTATACGG
168541	ATAGTTTGAA	TTTTACTCCT	TGATATTCCT	TTTATCATAG	ACATGCCACA	GTAGCTGGGC
168601	ACAGTGGTTC	ATGCCTCTAA	TCCCAGCATT	TTGGGAGGCT	GAGATGGGAG	GGAGACCAGG
168661	GGTTTGAGGC	CAGTATAAGC	AAGAAAGGCA	GACCATGTCT	CTACAAAAA	TAAAAAAATT
168721	ATCCAGGTAT	GGTGGGGCAT	CCCTGTAGTC	CTAGCTACTT	GGGAGGCTGA	GGTGGGAGGA
168781	TTGCTTGAGC	CCCAGAAGGT	TGAGGCTGCA	GTGAGCCGAG	ATTGCACCAT	TGTACTCCAA
168841	CCTGGGATAC	AGAGCAAGAC	CCTACCTCAG	AAAAAAAAA	AAAAAAAAA	AAAGTAGAGG
168901	TACCAGAGTG	ATATTTTCAA	TGTCACTGAC	CCTTCATTCC	CCAAATGAAA	ATCCCCCAAT
168961	AGGTGTTCAA	TTTTTACGTG	TCCTTCAGGA	GTTACTTCTA	AGATGAACCA	CTCTCTACCC
169021	TAAATGTCCC	TCCCCACCAC	CAAAACCAGG	GACCTCCAGG	CAGACATTTT	TGATGGTTTG
169081	TTTTCTTTAC	TAGACTGTAG	ATACCTAAAA	GGTGATGGGT	CTTTCTTCCC	TGTTTTCAGG
169141	CCCTACTGCA	TGGCTTTACA	TATTGTGGTT	TTTCAAATGA	TATTCATGGT	GTGAAACAAG
169201	AAAAAATGCG	GGTGTTTGGT	TTGAGAACAA	CCTGTTCTAA	AGCAAAAAGA	AATTCATCAT
169261	AACACAAATG	GATAGAGATA	AGAGTCCAAC	CATCCCATTG	AAGGTCAGGA	TGGACAGTCT
169321	AGATAATTGA	GCAAGAAATC	ATCATAAACT	ATTTTTCAGA	AGAATGACAT	GATGAAAGCT
169381	GTATTTCCAA	GTCATAATGT	TAGGTTTCAA	GTTAAATCAT	CTCAGCTCCT	GGGGAGCAGG
169441	ATAAGACTTG	GTACTTACCA	AAGCTCCCGG	GCCCACACAC	TCACCTTGTA	GCCCTGGCAT
169501	ACGTCTTCAA	CAAGAGCTGT	GGTGTGCCCT	TTGTGCTGTG	GTGCCCGCTC	ACAGCGCCAG
169561	CAGATGAGCT	GCCCCTCATC	TTCGCAGAAC	AGGTGGAACT	GCTCTCCGTG	TTCCTCACAT
169621	GACATTTCTT	GATCCGTCTC	TTTGAGGGCT	TCAATGAGGC	TTCCCAGCTG	CTTGTTGGGT
169681	CGGAGGCTAT	CCATATGAAA	TGGAGCCCGA	CACTGGGGAC	AGCAGAATGT	CTCCTGCCTC
169741	AGTTGCTTTT	GGCTTGGGTT	TTTAAAGAAG	TCTGTTATAC	ACAAGTGGCA	GTAGCTGTGT
169801		TGCTTACTGG				
169861	TCCATCATCT	TCTTGGTGCT	GGTGGTTGAG	GCCATAGCTT	TTATTGAAAA	GCTCCAATAT
169921	TGGCTCTAGA	GATGGAGATG	AAGCAGCCAG	AATTTTCCAC	CGTGATGAAA	ATACACCTCA
169981	CCTGCACCTC	TATGTGATGA	GCTGGCTGCA	ACTGACTTCC	ATAGGTCTTG	AAGGTTTTCC
170041	TTCCAACCCC	TATTATCTCA	TTTTGTATTG	AAGAAAAGAG	GACCTAAAAG	GAAGAAGTTG
170101	AGGCTGAGGT	TGTTTGGGCC	ACGTTTGAGA	ACTGCAACCC	AAGTGCAGAG	TTTCAAGTTG
170161	CCCTCATTAG	CAAGCAGTTA	CAAGTGGTTG	TTTAGAGGAA	AAAAAGCAGT	TTTAAAGCAG
170221	TTTTAAAGTT	GTTTGCCAAG	AATTTACATT	AAAATAGCAT	AAGCTTTTGA	CTGGCTATAC
170281		GTATTACAAA				
170341	GAACAAAATG	CTTTTAAACA	TGGGGTCTTA	ACTGAAGACC	TATACTCCTG	CCTCACTTGT
170401	CCTGATAAAT	TTTGCATACC	TCACATAGCT	CAGACTGCTC	TAAATTATTT	CATTATTTTT
170461		TCTTCTAACT				
170521		AGTGCAGTGA				
170581	GCGATTCTCC	TGCCTCAGCC	TCCCGAGTAG	TAGCTGGGTC	TACAGGTGTG	CACCACTACG
170641	CCCAGCTAAT	TTTTGTATTT	TTAGTAGAGA	TGGGGTTTCA	CCATGTTGGT	TGGCTAGGAT
170701	GGTCTCGATC	TCTCGACCTT	GTGATCCACC	CGCCTCAGCC	TCCCAAAGTG	CCAGGATTAC
170761	AGGCATGAGC	CACCGTGCCC	AGCCTCTTTT	TCTTTTCTTA	TAAGACAAGT	TCTCGCTCTC
170821	TTGCCCAGGC	TGTAGTGGAG	GGCAGTGGCA	TGACCACAGC	TCACTGCAGC	CTCGACCTCC
170881		CAATCCTCCT				
170941	CACCATGTCC	AGCTAAAGTC	TTCTCTCCAG	AAAGAAGAAA	TGCATTGGAA	TTTAGAGGAT
171001	ACACAAACAT	CTAGCTGTAT	AGCTAATACA	GTAGCCACTA	TCATGAGTAG	GAATTTAAAT
171061	TTAACTTAAT	AAAAATTAAA	ATGAAAAAAT	TCAGTTTTTC	TGTTCCAGTT	GCCACATTTT
171121	GATTGCTTAA	TAGTTGCATG	TGACTAGTGG	CTACATAACA	GCCTCAATAT	ACAACATTCT
171181	GTTATCACAG	AAAGTTACCT	TGGACCAAGT	GCTGGGAGAA	GCAATGCAGG	CTTCCTCACA
171241	AAAGCTGTAA	AAGAGAGAAC	TCAGGGAGTG	TGAAACTCTT	TCCTATTCTA	GTTAACTTCA
171301	AGAATAATTG	TTACCAGGCC	AGCACGGTGG	CTCACGCCTG	TAATCCTAGC	ACTTTGGGAA
171361	GCCGAGGCGG	GCAGATCACC	TGAGGTCAGG	AGTTTGAGAC	CAGCCTGACC	AACATGGCAA
171421	AACCTCATCT	CTACTAAAAA	TACAAAAAGT	TAGCTAGATG	TGGTGGTGCA	CACCTGTAAT
171481	CCCAGCTGCT	CAGGAGGCTG	AGGAAGGAGA	ATGACTTGAG	CTCCGGAGGG	GGAGGTTGCA
171541	GTGAGCCCAG	ATTACACCAC	TGCACTCCAG	CCTGGGTGAA	AGAGCGAGAA	TCTGTCTTAA
171601	АААААААА	AAAAGAATAA	TTGGTACCAG	AATTACTCTT	TGTAATTAGT	AGTAACACTT
171661	ATGCAATTGG	GTGATCTGTG	ACAGATTCCA	TTGAAGGAGT	ATGGGGAGCT	TCACCCCAAT
				_	_ _	

Figure 9 (Page 53 of 74)

WO 98/14466 PCT/US97/17658

142/162

171721	ATATGACTCC	CTGGTATAAT	GAGTATTTTG	AATTAAAGGC	CCTTAGAGAT	CAGCAGATGC
171781	TGGAAGAGAC	TTTTCCCCTA	TCTACATAAA	GACCAGTCAC	ACTAGACAAG	AAGAACAATT
171841	GITTTTCCTT	CCAACCCCTA	TTATCTCATT	TTGTACTGAA	GAAAAGAGGA	CTAAGAATGT
171901	AACCAGACCT	AATCAGACAC	TTTCACAAAA	TAATGTCTGT	CTCTCAGGCT	CATTCATTTT
171961	CCAAAGAGAA	CCATTTACAA	GTTAAACTCT	GTTCCTCCAT	TCATTCATCC	TCCCAAATAT
172021	TCATTTATTC	TCCCTAGTAA	TCATTTACTG	CCCCTCAAAG	AATTACCTAT	ATTCTCCTGA
172081	TATCACCCTT	CCCCTCTGAA	ATAAATATGT	ATACATGTAT	AAACGTTATA	CATACATATT
172141	TATACAGTAT	ACATACATAT	TTATACATAC	ATACATATGC	ATACATATTT	ATATTTATGT
172201	ATTTATACAT	AAGTATTTAT	AAATAAGGCT	ATATAAGTAT	CTACCCCCAT	TGGCAGAGGG
172261	GGTAATCACT	CTGTGATTCT	AGCCCATGTA	CTTGTTAATA	AATTTGTATG	CCTTTTCTCC
172321	AATTAGCCTG	CCTTTTGTGA	GTCGATTTTT	CAGTGAACTT	CAGAAGGCAA	AGGGGAAGTG
172381	TTCCCTTGGC	TCCTACACCA	TCATGACAAT	AAAATTTGAC	TCCACCTCGA	CCCCCCCAT
172441	CCCCCACAAA	GAACAACAAC	CAACACTGGT	TAATAAGGTC	GGTTGTTTTT	TGTTTGTGTT
172501	TTTGTTGTTG	TTGTTGTTGT	TGTTGTTTTT	GCTTTCAGGA	GCAGAGGTAT	AATAGGCAAA
172561	AGAAAGAGAA	AGGAGAATAG	TGAATACCTC	TTCTGCAGAG	AGGGGTGCCT	AAGTGGGACT
172621	TCCCTGGCTA	ATAACGTCTT	GCTAGAGACC	CAACCAGGAG	GATAATGGAA	GCAATCAAGG
172681	CAACCAGAAC	AACCAGAAGA	ACCAGTTTAT	CCTTTTTGTG	CCCTCTCCCT	AAACTGAGGG
172741	AATAAGAATT	GGAAAGAAGG	CTGCAGAGCA	GAGGGTTTGC	TCCTGAGGAG	CAGTTATTTC
172801	TATGGGATCA	GAGCTCCTGC	AGAACTGGGG	AGTTTACTTT	TACTATCTCT	TCTCCAGGAC
172861	AGGACCTATC	TCAAGAGACA	TGTTCAGAGT	GATTGCAACA	TAAAGAGTTT	GCAGACCCAA
172921				GAGGCCAGGG		
172981	CAGGAGCGAA	AAAGCCTGCC	TCTTCTGAGA	ACCTAGCTGG	GCTCTCCCTG	TACCCCCGAT
173041	CCCTCCCCCC	CGCCCGCCCC	CACACCCCTA	CTCCTGGGAG	CTCCTCTAGG	ACAGGGGCAG
173101				ATAAAAAACA		
173161				CAGTCTCTTG		
173221				CCTCCAGTGA		
173281	TGGGGTCACT	GCTCTTCTGG	GGAGATGGGG	CTCCCCTCCT	TCCAAGGCTC	CAGGGTTCCT
173341				TCTTCTGAGA		
173401				AAAACAAAGA		
173461				TTCCCCACCC		
173521				GACCTTTTCC		
173581				TGCTGCCACT		
173641				ACTGTGGGAG		
173701				TGCACTCCGC		
173761				AGCAATAAAA		
173821	CCTGTAAACT	ATCATGTGAC	CCCAACACAG	AGTATCTAAA	AACAGGAAGC	CTGCAGAGGT
173881				CTACTTTTGC		
173941				TTGCTATTTA		
174001				GGTGGTAACA		
174061	CACAAAATTC	ATCTGAGTCA	GCTTTCTATT	CTTCTCTGTC	CCGTTCTGTG	TCTTGTTTTT
174121	CTCCTTGCTG	TCCTTCTGCA	GGACTCAGAT	CTTCTTCAAT	AGCGAGGGTC	AGCCAGGATA
174181				GAGTGCCCCC		
174241				GTGGAGAAAA		
174301				ACTTCAAAAT		
174361				AAAAGATGGA		
174421	CTTGTCACAT	TTATAAGTCT	CAGGTGTAAG	AGGCATTTAT	GATAACAACA	TAATAAATGC
174481				ACCAGTAAGG		
174541				CACTGGCCTG		
174601				TTTGGCCCAG		
174661				ACGTTCCTTT		
174721	ACTCTTCAGC	ACTGCACCCT	CCTGGGTGCT	CACAGAGCCT	TCTGTTGTTT	TGCCACCTAC
174781	GATTCATCAT	GCCCTGGCAT	GATGGTTGCA	GACCCCATGC	ATAGCATGGG	ACATTCTACT
174841				AAAGAATGAG		
174901	CGATGAGTCC	TTGCAGATAT	CTACAACTTT	CATTGTTGTG	GATGTGACTC	TGTACCCAGG

Figure 9 (Page 54 of 74)

174961	CATGGCTCAT	TCCAGATCTG	TCCTATTGTC	AGAGGTGTTC	AAACCAGAAT	GACTCCATTT
175021	TGAATGGGGG	CTAGGTAAAA	TAAGGCTGAG	ACCTACTGGG	CTGCATTCCC	AGGAAGTTAG
175081	GCATTGTAAG	TCACAGGATG	AAATAGGCÅG	TTGGCACAAG	ACACAGGTCA	TAAAGATCTT
175141	GCTGATAAAA	CAGGTTGCAG	TAAAGAAGCT	GACCAAAACC	CACCAAAATC	AAGATGGCAA
175201	CAAGAGTGGC	CTCTAGTCAT	TCTCATTGCT	CATTATACAC	GAATTATAAT	GTGTTAGCAA
175261	GTTAGAAGGC	ATTCCCACCA	GCTCCATAGT	GGTTTATAAA	TACCATGGCG	ATGTCAGGAA
175321	GCTACCCTAT	ATAGTCTAAA	AAGGGGAGGA	ACGCTTGGTT	CTGGGAATTG	CCCACATCTT
175381	TCCCAGAAAA	CATATGAATA	ATCCACTCCT	TGTTTAGTAC	ATAATCAAGA	AATAACTCTA
175441	AGTATCTGTA	TTAGTCCATT	TTCACACTGC	TGATCCAGAC	ATACCTGAGA	CTCACTAATT
175501	TATACCAGGA	AAAAATGTTT	CATGCTCTTA	CAGTCCCACG	TGTCTGGGGA	CIGAGIAAII
175561	CCACAGCAGA	AGGCAAGGAG	GAGCAAGTCA	GGTCTTACAT	GGATGGCAGC	ACCOLACAG
175621	CTTGTGCAGG	GAAATTCCTT	CCTATAAAAC	CATCAGGTCT	CATGAAACTT	AGGCAAAGAG
175681	ATGAGAACAG	CAGTATAAAT	TACTCAGGGA	AAGACCTGCC	CCCATGATTC	ATTACCTCC
175741	CACCAGGTCC	CTCCCACAAT	ATGTGGGAAT	TTAAGATGAG	AGTTAGGTCC	CCDCDCDCC
175801	AAACCATATC	AGTATCCTTA	GTCCAGAAGC	TGATGCTCTG	CCTCTACACT	ACCONTRACTOR
175861	TTATTCCTTT	ACTTTCTTGC	TTTCACTTA	CTCTCTACAC	TTCCCCCAAA	AGCCATTCTT
175921	CACGAGATCT	AAGAACCTTC	TOTTAGGGTC	TECETTECES	CCCCCTTTCT	CONTROL
175981	TCAAAGGATC	AGGAAAAGGA	AGCTAGTGAA	TOCTARARAC	CARACARACT	GGTAACACTA
176041	ATAATAACAG	CAAGACAAAA	GCAAAACGCA	TTCTCACACAC	TCTCCCATCT	ACCATTACCA
176101	TCCCATTGCA	GGAAGGAGGG	GCTGGTTCNT	CCACACACTC	CCCDATTA	CACACCTGTT
176161	GGGGGTGCAG	ATGAGACTTC	AGGAATATGT	TCACAAACCC	ACCCCTACCC	GAAGCAGAGA
176221	CTGAACTATC	CCCAAGGAGG	AATGCATTAT	CTCTAATAGGC	TARACTTACC	AGAAATCAAC
176281	TGATTATGGG	ATATAGGAGT	CCAAAGACTC	DCDATCCOAR	CTACCTCACT	CTTGATCCTG
176341	TCAGAAGCTC	TGTACTGTGT	GTTCCCACTC	TCCCCAACAC	TCACCACTCA	AGAGICTCCT
176401	GAATGCCTTT	CCTCAACTCC	TTCACATTTT	CCCTCTCAAGAG	TARGUACICA	GCTATTCCTA
176461	GTTAGCAAGT	GTACCCCTCT	CTCCCTCCCA	A A CA TETTE CA	AACCCTATC	CTGACCACTT
176521	CACTTATCAC	TGAATATTTT	ACTA ATTTAT	WACATITICA	CTTTCCTTCC	GTTCCCATGG
176581		TGGATTTTTT				
176641		GTGATGTGCA				
176701	ATCACAAGAG	GTGTGTGCAC	ATATGTGGAT	CATCCACCTA	CIGAATIGCA	CTAAACACAC
176761	GGGGTATGTG	GTACTGTGTG	TECTETETET	CCTATCTCAT	ACATA CTTTTC	GGTGTTGTGT
176821	ATGCATGTGA	TGTGGTATGT	GTGTGCGTGT	CCATACATAT	TACCCCTCCC	GGGGAGGGGG
176881		TGGTACTAGA				
176941		CCCACCTGTA				
177001	GGCTGTGACC	TACTGGGCTG	ACCARATACIG	AIGGIIIGGA	CAGAGAAGAA	ATAAAAAGAA
177061	GAGTGGAGGG	GCCAAGGGAA	AGGAAATAAA	AACGAAAGTA	AAAGAAGAGC	TGGGAAAAGA
177121	AACTCACAAA	TTTATTAACA	TOTACACACA	CAGARAGAMA	GGAAACTTTG	CTGAAAAATC
177181		AAAGCTTATA				
177241	AAACTCTGTT	GATGGGATTA	CTCTTCCCC	GCAAAACAGA	TTATGGGAGG	GGAAGAAGAG
177301	TTTTTCTCTC	ACAGCCAGGA	ACAATTACCC	ATÉCACCCAT	CIRROLL	TAGGTCCGGG
177361	ATTTATTANG	TGAAAGGAAA	CCTCTCACCA	AIGCAGCCAI	CAAAGAATGA	GTGGAGTAGA
177421	TTTCCTCTTC	ACACTTCAAT	ACTACCCCTT	AAGACAAGGG	TECTGAAAGC	AGATTTCTGG
177481	TCCTACCACT	ACAGTTGAAT	ACTAGGGCTT	AAGACTCAAA	TTCCTGACAA	CTCCACCCTG
177541	TOCCATOTO	GCATGCAGGC	CITTAGACIG	AGCTACTCCA	TATTGATTAA	TTTCCTGAAC
177601	CAAGTTCCCT	TTAAGGAAAG TATCTGCACA	AAACATCCC	TOTALGOLOT	GTTTAGGCAA	GCCCCCTGTG
177661	CTCTGGGTAC	CATTCCCTTA	CTCTCTCCCT	IGIAAGCACT	TGTGGGGCAG	GTCAGAGGTT
177721	AGGGAGAGTA	AGTAGATCAC	CONTRACTOR	AAAGCAAGCT	GGCCAACTCC	TTTCATTACT
177781	TTCGGGCATC	AGTAGATCAG	TCCTCTTACA	CONNECTEDA	CATTATCTTG	TGAAAGTCCG
177841		GCATCTTAGG				
177901	GATAGTTGGT	GGATCTTAGG	TCACACACAC	AACTTTAATT	CCACGATGTG	TITTGGTAGG
177961	GGGCCATATA	GGCAGGGATG	A A TTTTCTCAC	CCCCAACAAC	ACCETT CAST TO	ATATGACCAA
178021		TTAGGGTATC				
178021		GTGAAAGCAA				
178141	ATCCTCCCAM	GTACAATGGT	ACGATCACAG	CICACTGTAA	TCTTGAACTG	GGTTCAAATG
0747	TICCICCAT	CIAAGCATTT	CAAAGTGTTG	GGATTACAGG	CATGAGCCAC	GGTACCCAGC

Figure 9 (Page 55 of 74)

178201	CTGAAACTG	C ACCCACTTTC	TGATAAACT	TTCAAATGAC	TAAACCCCAC	
178261		O DINGGAAGAA	L AUSUMACIALIANIC	א מייטי איזי איזייטי א		
178321		c coordiagi	- GCICACACC	י כדם מדריארא ר	*	
178381		T CIGORGGCCH	GUAGTTTTAL	· ACCAGGGGGG		
178441		_ ~~~~~~~~	ALL THE			
178501		" WATWARTET	AGCATTTTTT	كامل طحملهمليا بالماليان		
178561		A MOTICAGIGG	CGTGATCTCC	GCTCACTGCA	A C B C C C C C C C C C C C C C C C C C	
178621		- CCIGCCIICM	GCCTCCCAAC	TAGCTGGGAT	TRACE TO TO TO TO	~~
178681	CC11041	• • • • • • • • • • • • • • • • • • •	LITAATAGAG	ATGGGGTTTT	CCCATCOOK	aa
178741		- IGNCCICANG	IGATCTGCCC	ACCTTGGGGG	CTCCNTNOMO	
178801		CACIGCACCC	GGCAAAGTCT	' TAGCATTCTT	TACARAGE	
178861	***************************************	A GOGWGIWGIG	- AATTTCACCC	ייייי מידי ממממי		
178921		* WWWCICIING	AGATCAACAG	- ארברדא א ארבא	CACTEMBAGG	
178981	AAAATAGGA1	GGCCCCACCA	GCGAGAACAA	Tale tale to the tale to the tale tale tale tale tale tale tale tal	CTCCCTCCC	TAGGTACATA
179041	TGTGCATTAT	AGGAAAGACC	AAGAATGTAA	CCACACCTCA	CICCCICCT	GTTATCTCAT
179101	ATCAGTCTCT	AAGCATCATT	TAAATTCCAA	GGAGAACTAT	ACAGACCCTT.	TTATAAGATA
179161	TGATCCAATT	AGTCTCTCCT	GGTAGTTACA	TATTCCCCCT	CARCAGATTT	ATCTGTTCTT
179221	TGTTTCCCAT	AACCTATTTT	GCAAGGATCA	ACCCCCCT	CAACAGAATT	CCTCTTCTTC
179281	GCATATAAGC	TTCTAAATTC	CACTGGGATA	TTCCTACTA	ACTICITCAA	CTTCAAGTTG
179341	AGTAATTAAA	TTGTAAAGCC	TTTTTATCTTA	TIGGIACIAT	GTGCATGAGG	AGAACCACAG
179401	AGCAAAACTT	CCAAGGGCAA	ACCTATABAA	CAATCTGCC	TTTTTTTTTT	TTCATTTTTC
179461	TCTGAATAGA	CCAAGGGCAA	ACTORCOCCT	CAAAAATAAA	ATTCTAAAGC	CCCCCAACCA
179521	GCCATTAAGG	CTTTCTCTTC	TTCARCATCO	CTTAAAATGT	AACCTGAAAG	ACTGGCTCAG
179581	CACAGCTTTT	GAAGTGGGGG	ACAAACATGC	CTCATTATTC	CTCTCTGGCA	TTAACATCAA
179641	TAAAAACTTC	AAGTCTGATA	AGAAACATTT	TACAACCTAT	TCTCTCTGAA	GCCTGCTAGC
179701	TCCTTTCTAT	ATCCCATAGT	TCTTTATA CA	TCTTCACAAC	CTGTTATCAC	AACCTAGTGC
179761	ACTCCTCCGC	TAATCCCAAA	TCTTTATACA	AACTCAACCA	ATTGTCATCA	CCTCCACCCC
179821	ACGTATTTGA	TGCTTCCAGT	ATCCCTCCCT	CTCTGGACCA	AACCAGTGTA	CATTTCTTAA
179881	CACCTTGAGC	TTGATGTCCC	ACCACCTCCCT	AAAATGTATA	AAGCCAAGGT	GCATCCCAAC
179941	AATTTGGCTC	GCTTGTTCTC	TOTTON	GAGGGCTGTG	TCATGGGCCA	TGGTCACTCA
180001	AGATGACTGC	AGAATAAATC	CCTCCTCTCTC	TTTTACAGAG	TTTGGCTCTT	GTCATGACAC
180061	TCCCCGGATA	TTCACTGAAG	Character	AAGTGAGTGG	GGGTTTTGCA	AGGATAATTT
180121	GAACACTTCT	GCCCCAGAAG	LAGCIAGIAA	TAATACACTT	AAAGGTAGCT	AAAATGCATT
180181	ACTCCTGATT	TTTGTGCCAG	ACCTATGTCA	ACATTTGCTT	TGTGCCAGGC	TTATGCCAGT
180241	AAAACAGAAA	TGTTAATACA	TTCTAAATAA	AAATTCTGGA	GTTTCAAATA	TAATAACTGA
180301	TGGTGGCTCA	ATAAATAAAA	ATATATAATA	ACTGAAATAA	AAATTTACTA	AGGCTGGGGA
180361	CCAAGAGAGG	CTCACACCTG	TAATCCTGTT	ACCGGAAAGG	GGTCCGTCCA	GATCCAGACC
180421	GCA ACTITAT	GTTCTTGGAT	CTCACACAAG	AAAGAATTCG	GGCGAGTCTG	TAAAGTGAAA
180481	TGAGGGCTGC	TAAGAAAGTA	GAGGAATAAA	AGAACGGCTA	CTCCATAGGC	AGAGCAGCTC
180541	GGATAATTCA	TGGTCGCCCA	TTTTTATGGT	TATTTCTTGA	TTATGTGCTA	AACAAGGGGT
180601	ATTCCTAAAC	TGCCTCCATT	TTTTAGACCA	TATAAAGTAA	CTTCCTGACG	TTGCCATGGC
180661	TCATCCCCA	TGTCGTGGCG	CTGGTATGAG	CATAGCAGTG	AGGACGACCA	GAGGTCACTC
180721	ICMICGCCWI	CITGGATTG	GTGGGGAGCA	GTGAGGATGA	CCAGAGGTCA	CTCTCNTCCC
180781	TTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTT	TTGGTGGGGT	TTAGCCAGCT	TCTTTACTTT	TTTCCTTTTT	TTTTTTTTT
180841	TTCTCACTO	GCCCAGGCTG	GAGTGCAGTG	GCACGATCTC	AGCTCACTGA .	AACCTCCAAT
180901	CCCACCACAC	AAGCGATTCT	CGTGCCTCAG	CCTCCCAAGT	AGCTGGGATT .	ACAGGCATGT
180961	TA COCHEACAC	CCAGCTAATT	TTTTATATTT	TTAATAGAGA	CCGGGTTTCG	CCATGTTGCC
	CCCCTGATC	TCCAACTCCT	GCGCTCAAGC	CATCCAGCCA	CCTTAGCCTC	CCAAAGTGCT
181021	GGGCTTATAG	GTGTGAGCCA (CCCCACCTGG	CCTAGCCGGC '	TTCTTTACTG	CAACCTGTTT
181081	TATCAGCAAG	GTCTTTATGA (CCTGTATTTT	GTGCCCACTG (CCTGCCTCAT (CCTGTGGCTT
181141	ACAATGCCTA	ACTTACAGGG Z	AATGCAGCCC	AGCAGGACTC :	AGCCTTATTT (CACCCAGCTC
181201	CTATTCAAGA	TGGAGTCTTT (CTTGTTCAAA	TACCTCTGAC A	AAGCCCAACA (CTTTGGGAGG
181261	ATGACACAGG	AGGATTGCTT :	TAGCCTAGGA	GCTCAAGACC A	AGCCTGGGCA	ACACAGTGAG
181321	ACCCCATCTC	TAAAAAAAAA 1	AAATACAAAA	AAATTAGCCA (GCATGATGG '	гстстссстс
181381	TAGTCCCTGC	TACTCAGGAG	GCTGAAGTGG	GAAGATGGCT '	TCAGCCCAGG A	AATTCAAGGC

Figure 9 (Page 56 of 74)

181441	TCCATTCTC	~~~~~~~				
181501	NACCOTTO	GAGGCATTTG	AACCAGAATG	ACTCTATCTT	GAATAGGGGC	TGGATAAAAT
181561	AAGGCIGAGA	CCTGCTAGGC	TGCATTTCCA	GTATGTTAGG	CATTCTTAGT	CACAGGATGA
181621	GATAGGAAGT	CAGCACAAGG	TACACATCAC	AAAGACCTTG	CTGATAAAAT	AGGTTGTGGT
181681	AAAGAAGTTG	GCCAAAACCC	ATCAAAACCA	ACATGGCCAC	CAAAGGGACC	TCTGGTTGTC
	CARCAGGGG	ATTATATGTT	AATTATAATG	TATTAACATG	CTAAAAGACA	CTCCTACCAG
181741	CATCATGACA	GCTTACAAAT	ACTGCGGCAA	TATCTGGACT	TTACCTTATA	TGGTCTAAAA
181801	GGTGGAGGAA	CCCTCAATTT	TGGGAATTGT	CCACCCCTTT	TTTGGAATGC	TCATGAATAA
181861	TCCACCCCTT	GTTTAGCACA	TAATCCAGAA	ATAACTATAA	GTATGCTTAT	TTGAGCAGAC
181921	CACGCTGCTG	TTCTGCCTAC	AGAGTAGCCA	TTCTTTTATT	TCCTTACTTT	CTTAATAAAC
181981	CTGCTTTCAC	TTTACTGTAT	GGACTTGCCC	TAAATTCTTT	CTTGTGTGAG	ATCCAAGAAC
182041	CCTCTCTTGG	GGTCTGGATC	AAGACCCCTT	TCTGGTAACA	TCTTTCTGGT	GACCACGAAG
182101	GGACAATACT	GAGGAGACTC	TGAAGCCAAA	GGAAACAGAC	TACAGCACCA	ACTGGCTGAC
182161	TTTGGGTAAG	TGGTGGAGTC	CCCGGGTAAA	GGATAGGATT	GGGTTAGAGG	TGCAACTTAG
182221	GGGAGATAGG	GTCTCTCCTA	AGACAGAGAG	CGTTTCAGTC	CGCTCTTAAT	AAAGGGCAAG
182281	AATGCTTGAC	CGAACTTGGG	TTTGAGACCC	AACTTAGGAA	GGCTACAGTC	CTTAAGATTT
182341	AAGGGGTTAG	AGGCCCCTCT	CAGTAAAGTC	TCTCTTGGTT	AAAAACGGAT	TTAGCATTAG
182401	GGGATGTTAA	CTGCTATTCT	GTTTGTATTA	ATCTTCCCTG	TGCTCTTTGC	TGACAGCTAT
182461	GGGTGACAGG	ATTAGGCATG	TACAGGATCA	CGGGACATTG	GGAACTTTTC	TTCTCTCCAA
182521	AAGGGGAAGC	TTGACAGCTG	ATAGGACTGT	TGGAAAAGAT	CCCTTTGCTA	TGACAAGCAG
182581	CCGCCTGAAC	TTTTGATTCA	GTGTTGCTGC	AATGGGTGGG	TCTTTCTCTG	GCCTCTGTGA
182641	ACTCCTCACC	TTCCCCACCT	CACCACAGGC	AATGCTTTTC	TCCCTTTCTC	TCTTTTCTCT
182701	TTTCTGTCTT	TTCTGTTACT	TGAGACAACC	ATCTTGCCCA	GAGACCATAT	GTTGAAACTC
182761	CTGGTCAGAA	GTTTGATTAA	AGATGAAAGG	GCCTATCTGG	GGGCAAGTTT	GAGCCTTCCC
182821	AGTTAGATAT	TGGGTGCTAA	GTGGAGTGGC	CAATGTCTAT	GTTTTGTCAC	ATGTATATTG
182881	CTCTGGCTGA	AATGGAAAAC	GTTAATTTGG	TTACTTTATG	TGGCCATTGG	CCACCATCTT
182941	ACAAAAGTGA	GAGACATTTA	TTTGCCTGTG	GTTCCATGAA	ACAGAAAAA	GTTGGTTTTC
183001	CTTTGTGTCG	TAGCTTGGAC	CCAAGGGCTT	TGCAGTGAGC	AAGGTTGCTA	GCCCTCCTCA
183061	GTGAAAGAGA	ACCCAGAAAC	CTGGCATGCC	AGCAAAAGGG	TAAACATTTC	TTACCACTCA
183121	GGCTTCTGGC	CTCTCTCTCT	TAGTGAAAAC	TGAATGAATG	CTANANTCA	CTCTTTATCA
183181	CCTCTGTAAA	GTTTTGATTA	ATGGGAACAA	GGATTTGTGG	GGCTAGTCTT	DACCTCTAAT
183241	GAATCTGGTA	TACTTTGTGA	TATCAATTTC	TCTTTCTGTA	TTACTCTCTC	AAGCIGIAAI
183301	ATATGGTAGG	ATAGAACATG	GGCTTAGGAC	TCCATAAGCC	TOCTOTO	CCCACCCCAC
183361	TAAACTGGTC	CGTTGCAAAG	TTTATTACAG	GTCCCTGGAA	BABBABABA	TTAAAAAACTC
183421	GATGAAGTTT	CCTTCTCATC	TTGTTTTATG	TCCTTTGGAG	CTTCACCTTC	TANAAAACIG
183481	GCGGTACTTT	CTCTTCCTCT	CTGCCATCCA	GGGAACAGGA	ATTTTCCCCT	TAACCACGIG
183541	GTTAACTCTA	AAAATTATCT	CARCCATTC	CAAGCTCAAA	ATTTIGGGGT	TTATGTAATA
183601	TCTGGGAAGG	CCNATCCAAA	CTARCCATIG	CAAGCICAAA	ATTGGCTGCT	CTGGACCCCT
183661	TTATAATCCC	CCCCAACCTT	CIAACCAGIG	TTGTAGCTCA	GCAGCTAAGG	ATTTGTCATT
183721	TOTOTOROGO	TTA CCA TTT	CAATCCTGGC	TTAGGGAATG	AGTACTTTCT	GATTGATATC
183781	TTCCTCTCTC	TTACCATTTG	TTGATTCTGT	TCTCTTCCCC	TCCACACACT	GTCTTGAGTT
183841	CCCCTCTCTC	TGAGAACCTG	GGAGATTATC	TTTGGTAAAG	TTCAAAAGCC	AGAAATAATG
	GCCGIGIGGG	ATGGCTAAAG	TTGAGTAATA	AGAAACTTAA	AAGGACTCCT	TTTTTTTTTG
183901	CITTAGAGTG	CTATGGTTTA	TGGTTAAAAG	CTTAATTAAA	AGTGGATATT	CAATCTCTAA
183961	AAGCCTGGGA	CTCCTTGGGA	AAAGCAGAGG	AGGCACCACA	GACCCCATTT	TGGGAAAACC
184021	TCTGTTTTCC	TCATGAAACC	CCAGGAACTG	GAAGTGGATA	GATCCTTCGC	AAAATCTAAG
184081	GCTCTGTTTG	GCTTTGCATT	ATGTTATCTG	ATGTTTTTGA	CTTTTGGGGG	TATCAGAAAT
184141	TACTTTGCAT	TATGAGGGAG	ATCTGGTGTG	TAATAACCAG	GTAGGAAATA	TACTTCTGGG
184201	GATAGCTAAA	GGCAAATATA	GGTGAATACT	TGGCTATTTG	CACTTTTGGA	TCACAAGAAG
184261	CATTCTCTTG	ACTACCTAGA	AGGTATGGAA	ATGTCTCCAT	CCCCACCGAG	AGATAAGATT
184321	CCCAGGGGAG	ATGGCTGATC	CCCCAAAAGA	GGGCTGATTC	CCTCTTTTGG	GATCCAGGAT
184381	CTGGTATAAA	AATGGGACCC	TGGCCAGGCA	CAGTGGCTCA	CGCCTGTAAT	CTCAACACTT
184441	TGGGAAGCCT	CAGAGTTATG	AATGTCTCAC	CATACTGACA	CTTTGTGACT	GAGCTCCTCT
184501	CTACCCTGGA	CACAAGAGAC	CCTAATAATT	AGACAGGAAT	ATCATTGCCC	CTATTTAGTC
184561	TGAAGAAGTT	ATAGAAGATG	GATCTTTATC	CCACTGCAAT	CCTTAGGATT	AAGGGTTCCC
184621	TGGTAAAAGG	GAGTGGGAAA	ATATGTCAGA	GGCATTTGAA	TCAGAGTGAC	TCCATCTTGA

Figure 9 (Page 57 of 74)

184681	ATAGGGGCTG GGTABATAR GGGTGAGG
184741	THE TOTAL OF THE TANK THE TELEPOOR TOTAL T
184801	
184861	
184921	
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185041	
185101	
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185641	THE TAX TAXABLE ACTUAL CONTROL OF COMMENTS
185701	TOUR TOUR TOUR TOUR TOUR TOUR TOUR TOUR
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185941	TARGETT OF COMMENTER OF COMMENTS OF COMMEN
186001	TOTAL STANDAMAN ANNANT ACT BENEGGOOTHER AND CONTROL
186061	TICAGIGAAC ICAAAGATAG CTCAATTCAA ATCAMOMA OM OM
186121	
186181	THE THE RECEIPE GGACTCCTAG AAGGAGAAA GTCACACCAC
186241	AATGTTTGGA GAAATAATTT CTCAAAGCTT CCCATGTTTG GCAAAAAAAC ATTAACTTGC
186301	ATACATATTT TAGGAGCTCA ATGAATTCCA AGTAGGATAC ACTCAAAGAG ATCCATACCT
186361	AGACACATCA TAATCAGATT ATCAAAAGAT GAAGAAGATG AATCTTGAGA GCAGAAAGAA
186421	AGGAACAATT CATCACATAC AAATAGTACT CAAAAGATGT CTGGAGTAGG TATACTAATA
186481	TCAGACAAAA TAAACTTTAA GATAAGCATT GTTATAATAA ATAAAGAAAG GTATTTTGTA
186541	ATGATAAAAG TGTCAATTCA TCAAGAAAAC ATAACATTAT AAACATACAT GCACCTAACA
186601	ACAGAGCCCT AATATTCATG AAACAAAACT GACAGAATTG AAGGGAGAAA TAGAAAATTC
186661	GACAATAATA GTTGGAGACA TCAATACCTC ACTAGTTAGA CAAGATCAAC AAAAAAATAG
186721	AAGACTTAAC ACTTGAAAAC ACCTAACCTG ACCCTAACAT AAATCTATAG GTCACTACAC
186781	CCCAAAACAG CAGAATAAAC ATCCTTCTGA AGCTCACATG AAACATTTTT CAGGATAGAC
186841	TGTATATTAC TTCATGAAAT AACTCTCAAT AAACTCTCTTT CAGGATAGAC
186901	TGTATATTAC TTCATGAAAT AAGTCTCAAT AAATGTAAAA GGACTATAAT AATAGAGTAT
186961	ATATTCTCTG ACCAAAGTGG AATGAAGATA GAAATCAATA ACTAGGCTGG GCGTGATGGC
187021	TCACGCCTGT AATCCCAGCA CTTTGGGAGG CCAAGGCGGA CAGATCACGA GGTCAGGAGT TTGAGACCAG CCTGACCAAC ATCCTCAAAC
187081	TTGAGACCAG CCTGACCAAC ATGGTGAAAC CCTGTCTCTA CTAACAAAAT ACAAAAATTA
187141	GCCAGGCCTG GTGGCATCTG CCTGTAGTCC CAGCTACTCG GGACACTGAG GCAGGAGAAT CACTTGAACC CAGGAGGCAG ACATTGCAGT GAGCACTGAG GCAGGAGAAT
187201	CACTTGAACC CAGGAGGCAG AGATTGCAGT GAGCTGAGAT CGCGCCACTG CATTCCAGCC
187261	TGGGAGACAG AGCGAGACTC CATCTCAAAA TTAAAAAAAA AAAAGAAACT AGAAAAATAA
187321	GAACAAATCA AACCCAAAGC AAGCAAGAGG AAAATGAAAA ATTTCAAAGC AGCCAAGAAC
187381	AAAAGGCACA TTATGTACAG AAGAACAAGT GTATAGATCA CATATTTCTC ATAGACACAA
187441	TATAAGCAAA AAGACAGTGG AGCAAAATTT TTTAGATTAA TGAAAGACCT ACAATTCTGT
187501	ACCARGERAL ARACICCC CCAAATGAGG GTGAAATAAG ACAATTTAAT ACAGAGAATAA
187561	CAGGAAGGAA IIIAICTAGT CATATGTGAG AGTTTTATGA TACATTTTCT AGTGTATA
187621	TOGATOTTT CTATTE TARARASTCA ACCOTOCART TARATCOTAC ATTOTTCT
187681	TIGHTIIGA IIGACACAGI CATTAACTAA AATATTGTAG TATTTTTTA TOTGGCTGG
187741	TARAGGCAAT AAACATCTAA TCAGCAGACT AGAACAATAA AAAATATTTT TTAAAACTCC
187801	TILAGGUAGA ATGATAAAAG TCCCTTAGGC ATATTGAAAT TCCTATTTAT ACAAACCAAR
187861	AAACAGIACI AGAAATTGTA ACTATGTGAG TAAACAGATA ATATTTTTTC TCCATAAAA
,001	GTGGTTGACT ATTTTCACAA AAATAGTTAA CAATGTAATG TGTGATTTAT AGCATTTAAA

Figure 9 (Page 58 of 74)

DATE DESCRIPTION

147/162

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187921
         AGTAAAACAG GCCGGGCACA AAGGTTCGTG CCTGTAATCC CAGCACTTTT GGAGGCCGAG
         GCGTGCAGAT CACTTGAGGA CAGGAGTTCA AGACCAGCCT GGCTAACATG GCAAAACCCC
 187981
         ATCTCTACTA AAAATACAAA AATTAACCAG GCGTGGTGGT GCACGCCTGT AATCCCAGCT
 188041
 188101
         ACTCTGGAGG CTGAGGCACA AGAATCACTT GAATCCAGGA GGTGGAGGTT GCAGTGAGGC
 188161
         AAAATTATAC CACTGTGCTC CAGCCTAGGC AACAGAGCTA GACTCTGTCA CACACACACA
 188221
         CACACACAA AGAAAAGTGT ATGACAACAA CAGTGCAAAA GAAGCGGAAA TGAAAATAAT
 188281
          GTTATTTAT ATAAGTGGTA TACTTTTAGA TGAACTACGA TAAATTAATG ATGTATACTA
 188341
         TAAACTCTAA GGCAACCACT GAAATAATGA AACGAAGAAT TATGGCTAAC AAGCCACAAA
          AAGAAATAAA ATAGAATGAG AAAAAATATT TAAGTTGTTC AACAGATGGG AAAAAAAAGA
 188401
 188461
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         TAACTCTACC CATATAGATT ATCACACTTA AGGTAAATGA TCTAAATACT CTAATACAAA
188521
 188581
          AGCAGAGGTT GTCAGATTGA ATTAAAAAAA CAGACAACAA CAAAAAAAAG CAAAAAAAGA
         GCCACAACAT GCTGCCTACA AAAAATTCAC TTTAATATAA AGACACAAAT AGTCTAGAAC
 188641
 188701
         ACCATCACTT TTAACCTTAT TTACTCAAAC CTCCTAACTG ATCCCTATTT ATTTATTAT
 188761
          TTATTTATTT ATTTATTTAT TTATTTTGA GACAGAGTCT GACTCTGTTG CCCAGGCTGG
 188821
         AGTGCAGTGG CACCATCTAG GCTCACTGCA GCCTCTACCT CTCGGGTTCA AGCGATTCTC
         CTGCCTCAGG CCTCCCAAGT AGCTGGGACT ATAGCACATG CCACCATGCC CAGCTAATTA
 188881
          TTATATTTT AGTAGAGACG GGGTTTTGCC ATGTAGGCCA GGTTGGTCTC AAACGCCTGA
 188941
         CCTCAGCCTC CCAAAGTGCT GGGATTACAG GCGTGAGCCA CAGCACCCAG CTCCTCTTCA
189001
 189061
          TTTATTCTTG CTACGCTTCC TCCAATCCAT TTTGTGCATT TGATGATTTT GCCAGTAACT
 189121
         TCTTTATTTT TCTGGTAAAA TTACTTATGG GTCACTGAGG ACTGGGATGT TCTTTCTTCT
          AGAGGGGGTT TGTGTCTGCT TTTGCCAGGA AGCTGGGGTA CCACCAGTCA AGTATTACTT
 189181
 189241
          TAAACTCAAT TCATGAATTG AGACTTTTTT TTTTTTTTT TTTTTTACGC AGAGTCCTAC
         TCTGTCACCC AGGCTGGAGT GCAGCGGTGT GAACATGGCT CACTGCAGCC TCAACCTACT
 189301
         GAGCTCAAGC AATCCTTCTG CCTCACCATT CTGTATAGCT AGGACTACAG GTGTGTGCCA
 189361
         CCATGCCTGA CTAATTTTTT AAATGTTTTT TTTAGAGATG GGGCTCACTT TGTTGCCCAG
 189421
 189481
         GCCGGTCTCG AGCTCCTGGG CTCAAGTGAT CCTCCCACCT TGGTCTCCCA AAGTGCTGGG
 189541
          GTTACAGGCA TGAGCCTCTG TGGCTAGCCA AGACTTTTTA TTTTTTAGCC TAAATGTGTA
 189601
          TAAAAGTTGG CTTGTGGTTA CAACTTATCA GGATTGATGA TCTCTCTCTC TCTCTCTCT
 189661
          TCTGTCTCTC CCCACCTCTC TCACATCCCT TGCTCTGCTG AGAAGCAGAG CAAACATTCT
          AGCAGTTTCC AGAGAGTAGG ATGGGATTAC TTCTAGTTTA CTTTTATCAT CCTTTGGGAT
 189721
          CGCAGTATTA CTGGGAGAAC ACAAGTATCT CTTATTAGAC ATACCACCTT TGTAGAATCT
189781
          GGACTTTCAT TTTAGACTTT ATTTGTTTTC TACTATAAGC AATTTAAGTT ACAGATCTCT
 189841
          CTACACACTG TTTAAGTTGC ATCCCATGAA TTTTGATGTG CTTTATTGTC ATTATTATAT
 189901
 189961
          190021
          TGTTTAATTT CCAAATATGT GTGTTTTTTT CCTACATTTC TTATTTTTAT TGATTTCAAA
          TTTATTTCTA CTGTAGTCAG ATTTAATAAT TCATTTATTT TTATTATTTT CATTTTTTA
, 190081
 190141
          GAGACAGGGC CTTTCTGTGT TGCCCAGGTT TGTCCCAAAC TCCTAGTCCC AAGCAGTTCT
 190201
         CCTGCCTCAG CCACCCAAAG TGCTGGGATT ATAGGCACGA GCCACCCGTG CACAACCAAC
 190261
          AATTCATTTA AAAAGTGGGC AAGTGAACTG AACAGACATT TCTCAAAAGA AGGCATACAA
 190321
          TTGGCCAACA AATATATGAA AGAATGCTCA ACATCACTGT ATTAGTCTGT TTTCATGCTG
 190381
          CTAATAAGA CTTAACCTGA GACTGGGGAA TTTACAAGAG AAAGAGGTTT AATGGACTTA
 190441
          CAGTTCCACA TGGCTGGAGA GATCTCACAA TCATGGTGGA AGGCAAGGAG GAGCAAGTCA
          CATCTTACAT GGATGGCAGC AGGCAAAGAG AGAGCTTGTG CAGGGAAACT CCCGTTTTTA
 190501
          AAACCATCAG ATCTCGTGAG ACTCATTCAC TATCATAAGA ACAGCATAGG AAAGACCCGG
 190561
          CCCATAATTC AGTCACCTCC CACTGGGTTC CTCCCAGGAC ACATGGGAAT TGTGGGAGTT
 190621
 190681
          ACAATTCAAG ATGAGATTTG GGTAGGGACA CAGCCAAACC ATATAAATAA CTAATCATCA
 190741
          GGGAAATGCA AATCAAAACC ACAATAAGGT ATCATCTCAC CCCAGTTAGA ATGGCTATTG
 190801
          TCAAAAAAC AAAAAATAAC AAATGCTGGT GAGGATGTAC AGAAGAGGGG ACTCTTATAT
 190861
          CCTACTGGTG GAAATGTCAA TTAGCATAGC CATTATGCAA AATAGTATGG AAGTGAGGTA
 190921
          GGTTACATAG GGTGGTCACA GCCTCCCTTG AAAGGAAACA AGAAACTTGT CAAATTGATG
 190981
          GAGAGAACAA ATCTCTTGAC ATTACACAAA CTGCATCTGG GGCTAGTGGT TAGAATATCC
 191041
          TCAGTCAAGG AGGTAGAAGA GCAGGAGGGA AAATCCCTAA GTTCGTGCAA GTGCAGAAAC
          CCACAAGCTG TGTTCTCAGG TTGACATATA CTCATTTTAA TAGTAAGAAA CACACCCTTG
 191101
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Figure 9 (Page 59 of 74)

191161	GGTAGAGAAT	TAAAATGCTA	ATAATACATG	TGATGTATGT	ACTAGCGTGT	ATGGCAATAT
191221	IGCAIGCACA	TTCAAGAGAC	CACCCAAAAC	ATATTTAACA	ACAATGCCCA	TTCCCACCCC
191281	CICAIGGATA	ATCACGTAGG	ACTCCCATAA	CGGGAGTTTC	TTCAGTGTCA	ATTGGTGCTC
191341	AAGTAGCCGA	CCCTGACTCT	' GCTATCAGCG	TGTACTTTCA	CCTTGCAATA	y y Car Commence
191401	CCTACTTTA	CTTTGGACTG	GCTTTCAAAT	TCTTTTGTGC	AGGGAATTCA	AGAATCTCAA
191461	CCAGCCCACT	GACAACAGAG	GTTTCTCAGA	AACCTAAAA	TAGATCTACC	AGATGAGGCT
191521	GAAAATCTGC	TACTGGCTAT	TTATCCAAAG	GGAAGGAAAT	CAGTATACA	AGATGAGGCT
191581	ACATCCCCAT	GTTTATTGCG	TCACTCTTCA	CAAGAGCTGA	TATATAGACT	CAACCCTAAA
191641	TGTTCATTAA	CAGACAAATG	GATAGAAAAT	GTGGCATATA	TACACAATGA	LAACCCTAAA
191701	GGCCATGAGA	AGAATGCAAT	CTTGTCATTT	GTGGCAACGT	AGATGAAACT	CCACAACTATTT
191761	ATGTTAAGTA	AGATAAGCTA	GGATTGGAAA	GATAAATACT	ACATGTTATC	AGRICATI
191821	GAAAGTAGAG	AAAAATTTTT	AGCTCATGGA	TTTAGAGAAC	AGAACTGTGG	ACTUATATGT
191881	CTGGGAAGGG	TAGCAAGGAG	GGGAGGATAG	GGAGAGGTTG	GTTAATGGTG	ACARAG
191941	AGCTAGATTG	TAGAAATGAG	TTCCGGTGTT	CTGCACCATT	GTAGGGTGCA	TATOCTTA
192001	TCTCATTTAT	TGTATATTTT	CAAAAAGCTA	GAAAAGAATT	TTGAATACTC	ATGGTTAAC
192061	AAATGATAAA	TGTTTAAGGT	GATGGATATA	CTAATTACTC	TGATTTGATT	ACAACAAAAT
192121	GTGTACACAT	ATAAAAATAT	CACTCTTTAT	CCCGTATATA	TGTACAGTTA	ATTACACATT
192181	ACTAAAAATA	AAAGAAAAA	AGAATATGAT	CTATCATCAT	GTATATATCA	TTATATGTCA
192241	AGCAAAATGT	GCATGCAGAT	ATTGTGTATA	ATCTTCTATA	AATCAATTAG	TGTGTACTTG
192301	TAGATAGGAT	TGTTCAGATC	TTCTGTGTCT	TTACTCATA	TTTGTCTAGT	CTCAAGATAA
192361	TTACCAAAAA	AAGGGTGTTA	AACTCTCCAA	ATCTCATTCT	AGAATTGTCT	TATTGCATCA
192421	TTCTTTTCCA	TTTTTACTTT	ATGTATTTTG	AAACTCTCTT	ATGACATTTT	ATTTTGTCTT
192481	TTAAAACTTC	GTTATGTATT	TTGAAACTCT	CTTCTTACAA	TCATACATTT	GCTATGTATT
192541	TGTTTTCTTG	ATGAAATGAC	CCTTTTCTAT	TOTCOTTOTT	TTTGTTTTTT	ATGATTATTA
192601	GTCTCACTCT	GTTGCCCAGG	CTGGAGTACA	GTGGGAGAAT	CTTGGTTCAC	CTGAAATGGA
192661	ACCTCCTGGG	TTCAAGCGAG	TOTOCTCACT	CACCCTCCAA	GTAGCTGGGA	TGCAACCTCC
192721	GTGCCAGCAT	GCCAAACTAA	TOTOCIONCI	TTDTTTTTTT	CAGAGTTTCA	Tracaggeat
192781	CAGGCTGGTC	TCGAACTAA	CACCTCACCT	TIATTAGAGA	CAGAGTTTCA	CCACGTTGGC
192841	TTTATTTTT	TERGREDER	TOTOLOTO	GATCCGCCCA	TAGAATGCGG	TTTATTTTAT
192901	TTGGCTCACT	GCAACCTCCG	CCTCCTCCCT	TCACCCAGGG	CCCATGCCTC	TGGTGTGATC
192961	GTAGCTGGGA	TTACAGGCAC	ATCCCACCAT	CARGCAATT	TTTTTGTATT	AGCCTCCCGA
193021	ATGGGGTTTT	TCTATGTTGG	CCACCACCAI	AACTCACTCC	TTTAACAATA	TTTAGTAGAG
193081	CTCTGTCTCT	GGTAACACTC	TCTCTCTTA N	AACTGACTCC	AGCTGTTATT	CAAAATATCA
193141	TTTTAGTCTT	TTTATCCTT	CTCTTTCCAT	ACTOTATITI	TTTTAATATG	ATTATAGCCA
193201	AGTTATCTGT	CTTTTTTTTTTTT	TTANCATCTT	AGIGIATATA	CCAACGTGTT	TTTATTCTCA
193261	ATTTTTAAGT	CGATTCTAAC	AACCTTTCCC	TCTCTTCTAG	AATATTTACA	TGGTTCTTGC
193321	CTAACATTAA	CONTICIANC	TOTTTOCK	COLCACTICA	AATATTTACA	CCATTAACAT
193381	TGAACATAAA	GTGTGATAAC	TCLLLCCACA	GTACACTGGC	TAGCATCTCC	CATATAATAT
193441	GGGGTGGAGA	AAGCATTCAA	CAACAICCIT	ATTTCATTCC	TACTCTGAGT	GGAAAGGGCA
193501	CTTCTGCATT	AAAAAAAAA	ATTACATTTE	CCAMCAAMMA	TCTTTTTGTT TTAGGAGAAA	ACACTGTTTT
193561	ATTTTCCTGG	AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA	ATTACATTTT	GCATGAATTA	TTAGGAGAAA	ATATTTTCCA
193621	TTTACATAAC	CTACATAACA	CACACGICI	CTCAATTTTG	TTTCCATCTT	TCTTCCACAT
193681	TTCTCTCTCT	CINCAINAGA	TTRECALLAR	CAAGTATATT	TTACATGGCT	TCTCAGTGTC
193741	CCATATCGTT	TTCCCTTTT	CACACCCTA	ATGGCACTCT	TGTATTTCTG	GTGGCTATGT
193801	ACATTTCATC	TCCTTCTTCT	CALMOL MORE	CTACTTCTTT	CACCAGTATT	AAAGACATGT
193861	CCTAAACACT	CCATTA	GGATGATTTT	AAATGACTCA	AGCTAATAAT	CCTAATTTTA
193921	ATTACCCTCC	CCATTATTTT	AAAATGTATT	CCTTTATGCC	CACAATAAAC	ATTTATTGAC
193921	TATTCACCC	ACAITAGGCT	TCTCTATGGC	AGACATTAGG	CTGGACCCTA	GCCATATATC
	TATIGAGGGA	AAAAAAATTA	TITICTATAT	AAGTTTCCAG	AAAGCCAAGA	TGTGTTTTAA
194041					TAAGAAAAAG	
194101	GAGAGAAGAA	CAAAGCAGCA	AGCAACTCCT	GGAAGGACCA	CTGCTGCAGA	GGTAATAACT
194161 194221	GGTGAACCAT	GTTTTGGAGA	AGGAAAAGGT	CACCAAGAGA	AGGAGGGGT	CCAGGGTGTT
					AATTCCGTAT	
194281	TGAAGTTCCA	GGACCATGAG	CTTGGAGAGC	ATGAAGTACA	GGAGGAGGGT	TGGTTTCAAA
194341	TAAATCTGGG	AATGAAACAG	TGAAGCCTCT	GGCAGAACTC	ACATCTCTTT	CCTCCCCTCT

Figure 9 (Page 60 of 74)

194401	TCCTTGCACA	TTCCCTTTAT	GGAGTAATTG	CAGGGATGGG	AAAAGTTCAA	A A C C A C C A C T
194461	GAGCCTAGGA	AGTGCTAGGG	TAAAGTGGAG	AATGAACCTG	CGTGATTTCC	TCATCCTAAA
194521	CTAGGTTCTT	CTAGGAGAGC	CCTTCCCCAT	AAAATCTGCC	CTCCTCCAAC	CCCCCCACAC
194581	AGCCTAAGCT	CACCTCCCAA	AGACCCCTTA	CTTGCTGACT	GAATCTCATT	CCACCCACAC
194641	ATGGCCTAAA	ACCCTTCCAT	AACTCTATAG	CCAAATTCAA	TTTTACACAC	CCACCCAGAC
194701	AACCTTTCTT	CCTCTAAGTC	TGCCACCCTA	GGCAATTCTC	AACATTCTCT	ACACACTERC
194761	GGGCCATAGA	CGTGCTACCA	AGTCTCCAGA	CCTAGACCTG	ATGGAGCACT	CCTCTAATCA
194821	GACGACCACT	GGCCTTTGAA	CCAGACCCTT	CTCTGTGGCT	CCTATCCATC	GCIGIAATGA
194881	TTTGAGCACT	GCTGCCAAGA	CATCTTTGGC	ACTTTGTTGT	COLNIGCAIC	1 CCAACCTGT
194941	ATCTACAAAA	CACCTAACCT	TTAAAAATTC	ATTGTCATTT	CATATCATCA	AACIGAACTA
195001	AAGGCCAGGA	AACTGTTCCA	GGTTAATAGA	GACTAAAGAG	ATAGCARCA	AAGATAAAGA
195061	GTGATCCTGG	ATTGAGGGGA	AAAAGTGTTG	TCAGAGACAT	GATTGGGACA	COMCOMINA
195121	TTTGAATTTG	AATTTAAAGA	TAAAGTATTG	AGTAATATAG	GARGATCATT	DECEGRAAA
195181	TTCAAATGTT	TCAGTAAGTA	TATATATATA	TAAAGAGATA	TABAGACATA	AICIGCAACT
195241	GATGGATAGG	TAGAGAAAA	GCAAATGTAT	AATATTAACA	ATCTACCTAA	TAAATAAATA
195301	AGTGTTCTTT	GTACTGTTTT	TCTGATTTTT	CTATATGTTT	CAAATCATT	AAAGTATATG
195361	AGGTTTTTGG	GGTTTTTTTG	TTTGTTTTTT	GTTTTTAGAG	ACACCATCAT	1 AAAA TAAGA
195421	CCAGGCTGTA	GCTCAGTGGC	CCAATCATTG	CTCACTGCAG	CCTCAACTTC	ATTCTGTCAC
195481	GTAATTCCCC	CTACCTCAGG	CTCATGAGTA	GCTGGTACTT	CACCTCTCCA	CTGGGCTCCA
195541	CAGCTAATTT	TTATTTTTA	AATTTTTGTA	GAGATGGCAT	CAGGIGIGCA	CCACTGCACT
195601	AGTCTCAAAC	TCCTGCCCCC	AAGTGATCCT	CCCACTTTGG	CCTCCCAAAC	CACCCAGGCT
195661	ATAGGCATGA	GCCACTGCAC	CCAGCCCCAA	ATAAAAAAGT	ATTTTATTT	TGCTAGAATT
195721	TTAATTTTGA	GTCAGAGTTT	CACCCTTGTC	ACCCAGGCTG	CACTCCAATC	AATTAACTAA
195781	GGCTCACTGC	AAACTCTGCC	TCCTGTGTTT	AAGCGATTCT	CTTCCCTCAC	CATGATGTT
195841	AGCTGAGATT	ACAGGTGCCT	GCCACCATGC	CCAGCTAATT	TTTATATAT	ACTOCTGAGT
195901	GGGGTTTCAG	CATGTTGGTC	AAGCTTGTCT	CAAACTCCTG	ACCTCACCTC	TAGTAGAGAC
195961	CTCGGCCTCC	GAAAGTGTTG	ATGAGCCACC	ACACCCGGTC	TARARACTAT	ATCCACCCAC
196021	CAGTCCCACT	CTACCTTGTC	CTACACTACC	AGGGGCTAGG	ATCACCCCAT	TTTAAAACCA
196081	CTATGAGATA	GAGGAATCCA	AGGAAGAAGA	TAAGCTACTT	CCTTCCTCT	GICTICTAGG
196141	TGTGTGCTCT	CATGTGCTCT	CTCTCTCTCT	CTCTCTCTCA	CACACACACA	TAGGGTCTTG
196201	CACACACACA	CACACACATG	AATACCAGAG	CTATCACTTT	CCCACACACA	TACACACACA
196261	ATCCCAAGGG	TTTTGTGTTG	TAGTGGTTTG	CTCATTTGTT	TCTTTTCTTT	CTTTCCTCTC
196321	ATTATTCTTT	TTCTCTTTTT	GCAGCTGAAG	GGAGAATTTC	CAGGCCAGGC	CTTTCCCCC
196381	TAGAGTTACA	GTGCCTCTAT	TCAGGCTTCA	TAGAGAGACC	TEGENTERA	TRETEGECCAT
196441	CTTTTATCCA	GTTCAAAATA	ATGCATTCTC	ACCAAGATGT	ACTTTCARA	TAGTGGGGG
196501	TAAAACACAA	AATTTTATTT	ATGCTGAACA	TTGAATCACT	TTTTTCTCTCT	AAAACAATAC
196561	AAAGTTATAC	ACACACAAAC	ACATTTGCTC	CTGCTTTGTT	TATTCCCCCA	TTTTGTGTAG
196621	TGGTAATACT	TCATCAGGCA	TGAGTAGTAC	GTCTTGGAAG	CTCTCCTCTA	GGGGTATGTT
196681	TCCTATCTGC	TTCCTTCAGC	ATTCTCCAGT	GTATCTGTCA	TCTCTCTACC	AAGCCTAGAC
196741	GTCTCCAGAA	CTTCCATTCA	CATTTAGAAG	AGGGCAGCGG	CTTTCTATCC	AAAATATA
196801	CTCTCATTCA	TCTCTATTCC	TTCTTCTAGC	TATGGTCCAG	CTCACCTCTT	TCCDDTDDD
196861	TATCTATATG	AAGTCTGCGA	ATGGTTCTCA	GACTGGTTGA	ACATTAGAAT	CACCECACEA
196921	CCTTCTAAAA	TTCTTATTAC	CCAGGGCATA	TCTCAGAATG	ACTACCACAC	CACCIGAGIA
196981	GGATTAGGGA	TCATGATCTC	TGGAGTCTGG	TTTAGGCACT	AGTGCTGTTT	A B B B COTA COT
197041	TCATGAGGTG	GAGGTTGCAG	TGAGCCGAGA	TGGCGCCACT	CCACTCCAAC	CTCCCCCA CA
197101	GAGTGAGAGT	CTGTCTCAAC	AACACAAAAC	AAAAAAAACC	A A CTA CCCTT	CTCATTTCAA
197161	TGTCCATCCA	AAATTGAGAA	CCATTAGGTA	AGGCCAAGCT	GTATAATTAA	ACACCACTTT
197221	TCATTTGTCT	GGTGTGGTGG	CAGCTTTTTG	ATAAGGGAAG	TATTCTTCC	ATCCACATAC
197281	CTGAGCCTCA	CTCCTGAGAA	CACTGGTGTG	TATGTTGCTA	ANTTOTIGCC	GGTGATTCTC
197341	AGGTTCCTTC	CTGGATAAAA	ACCACTGACC	CTGGGAATGT	ACCCACTGCC	AATCTCCTCC
197401	GTAAACCTTG	GATACTGGGA	AGCCTACAGT	TGAAAATATT	GGGCTTGAGA	TCCTCANACA
197461	AATCTTGTAT	TTCATTAAGA	CTAATATTTG	GTACAGTGCA	GCAAATCAAG	GGAATTTTGC
197521	TGGCTGAGTT	CTTTTAGAAC	TTTTGCATTG	AAATAGGTTC	ABCCBCCAAT	DAGTTAAAAC
197581	TACAACCTCA	GCTAAAGGAT	TAAAAGACAC	GTGAGCTGGG	TAGGATCACC	TCTNNCNTTC
					TAGGATGAGG	ICTAAGATIG

Figure 9 (Page 61 of 74)

197641	CCTCTCCCCC	CTCNTN CCTC	#11			
	TCACCTORORCO	CICATACCIG	TAATCCCAGC	ACTITGGGAG	ACTGAGGTGG	GTGGATCACT
197701	TOAGGICAGG	AGTTCAAAAC	CAGCCTGGCC	AACATGGTGA	AAACCCATCT	CTACTAAGAA
197761	CACAAAAAA	TTAGCTGGGC	GAGGTGCCAG	GCACCTGTAA	TCCCAGCTAC	TGGGGAGGCT
197821	GAGGGAGGAC	AATCACTTGA	ACTCAGGAGG	CAGAGGTTGT	AGTGAGCTGA	GATCGCACCA
197881	CTGCACTCCA	GCCTGGGTGA	CAGAGCAAGA	CTCCATTTAA	AAAAATAATA	ATAATAATAA
197941	CAATAATAAT	AATTCAGACA	TATCCAGGCA	TCAAACAGAT	ACCTGGGGCA	GATGAATAGT
198001	CTTGAGATTC	AAGTCACACA	TGAAATTTAG	GTGGAAAATG	ACATTGGAGA	AATTTGAGAT
198061	TATGATGAAT	GGAAATTTTT	CAAAGAGGAA	TTTCAGGCTC	TGTTCTTGAG	GGGATAGATG
198121	GACTTCCAAC	AGCAATAACA	CAGGATTAAT	GAGGACTTGG	GATGTTACAT	AAATTAGAGA
198181	TGTTAGATGG	ATAAAGAGAT	AAAAGTACTC	TCTCTAAGAA	CATGGGACCA	GAGATAGGCT
198241					TAAAAATAAA	
198301					TCAGTTTTTC	
198361	TCTTTAACTA	CAGGGTTGGT	CTGGGTGTGC	AACACAAGAA	AGCCTGGCAT	ATACATGGAT
198421					TTCATGTATT	
198481	TCTGTTTTTT	CCTTCATTGA	AGTCAATGGC	TGATATTAGA	TTCTACTATT	CATGTGTACT
198541	AGTTATATAT	AATTGTTACA	AAACAAATTA	GCAAAAACTT	AGTGGCTTAA	AGCAACACAC
198601	ATTTATTATT	ACCTAAGGTC	TGTGGATAGA	AGTTCTGACA	TGGCTTAACT	GGGTTCCCTG
198661					TCTGAATTCT	
198721					GTTGAAAAAT	
198781	GCACCGGTAG	AAGCTTCTTG	GTAGAGGCTG	ATTCAACTTC	TAGAGGCTGT	CTGCAGTTCC
198841	TGTCACCCAG	GGTGGAGTGC	AGTGGAGCAA	TCATAGCTCA	CTGCAGCCTT	GACCTCCCAG
198901					GACCACAAGT	
198961					GAGAACTTTG	
199021					AGCCTAAAAG	
199081					AGCAGGACAA	
199141					ATGGTCCTCT	
199201					TTTTCAACAG	
199261					CTTTTCTTGG	
199321					CCTAAGATGA	
199381					GCCTCCGTTC	
199441					AGACCAGATC	
199501					TAATGAGTGT	
199561					CAACTTTTGG	
199621					TTCTTTTCAT	
199681					CAGAGAAATA	
199741					ATTCCCTCTT	
199801					CCCCACGTT	
199861					CCATCAGACA	
199921					TGAAAACATT	
199981					AGAGTTGGTC	
200041					CACGTGCATG	
200101					TCTAAAATTT	
200161					AGTCAAATTT	
200221					CTTCAGAAAA	
200281					TTCTGCGTCC	
200341					TCCTGGACAT	
200401					GCTATTTCCT	
200461					GGTTTTTGCC	
200521					TACAAAAAA	
200521					ACTACCTTAG	
200581					GGGGTGTGTG	
200701						
200761					ACAACGCATC	
200761					TTGAAAACAT	
400041	CIIGIATATA	IACACACACA	TACACATACA	IGCAIGTATG	TACATATACA	CATACAGACA

Figure 9 (Page 62 of 74)

200881	AAAATGTATC	CTATGTATAT	TCACACATGT	ATACACACTC	ACACCTA CAM	101000000
200941	ATCCATAGTT	TATAAATGTT	CCTTTTTTTT	GGTCACCTTT	TTCCTAACAT	AGAGTTTTAC
201001	TTTTTTTTT	TTGAGACGGA	GTTTTGTTGT	CATTGCCCAG	CCTTACTCC	TTACACTTTT
201061	CTCACCTCAC	TGCAACCTCG	ACCTCCCCC	TTCAAGCGGT	TOTOGRAGO	GTAGCGCGAT
201121	AGTAGCTGGT	ACTACAGGTG	TGCGCCACCA	TGCCTGGCTA	ATTTTTTTTT	TAGCCTCCTG
201181	GAGACGAGGT	TTCACCATCT	TCCCCAACCA	GGTCTGGAAC	ATTITITITAG	TITITITATA
201241	CCTGCCTCAC	ATTCCCAAAC	TCCTCCCAAGCI	GGTCTGGAAC	TCCTGACCTC	AAGTGATCTG
201301	CTTACACATC	TTTTTTTTTT	IGCIGGGAIT	ACAGATGTGA	GCCACTGCAC	CCGGCCAAGT
201361	CCTACACATC	CTCLCLCL	CACTAAACTG	TTTACCCAAA	CCTGATAACC	CAAGTCAACA
201301	CIAIIAIGG	CICACACAAT	CITATGTAAA	CAAAGATACA	GATATATAGA	ATTTTCTTGA
201421	AAAAAAAAAAA	TA CAMMAMAIG	GAGTCCCTTT	ATACGTCCTT	AGTATCTGCT	TTACTCATTT
201541	TTTTA	TACATTATAT	GAAAGTATTC	AGGTCAAATG	TTATAGATGT	GATTCATTCT
201541	TITIAACIGI	GITATTITIC	TGCAATGACT	ATGTATCACA	AAGTACTCAG	TCTTCCACTG
	AIGAAAATTT	GGGCTATTTC	CAGTTTGTCT	TCCATTTTTC	TTTCTTCCTC	TTGGATTTTC
201661	ACTCAATGTG	TTTACTAATT	TAGGAAGAAT	CAATAGTTTT	TATGGTATTA	CTTCTCCCAT
201721	TCAAGAATAT	AGCATATGGT	ATAGTATAGT	AGAGTACTTA	GTTTAATTTA	GCCAGATCCT
201781	GTTTTCTGCC	CTTTAATAAA	ATTCTATCAT	TTTCTGCCTT	TGAGTCACAT	TTTCCTTGTT
201841	CATATAATTC	TTAAAAAAATG	TATAGTTTTC	ATTCTAAGGG	AACATAAAAA	CTTCTTTCCA
201901				TGGGAAAAGT		
201961	ATCTTTCCAG	TCAGTTCACC	ACATTTCCTT	TATACCTTTG	TACTTTAATC	CCCAGTCATG
202021	TTGAACACTT	CTTATTCCTC	ACACCAAGCC	TCAACGGGTT	TGCTCTTTCT	GGAAGGTGCT
202081	TCCCCTGTAT	TACTGACTTA	TTCATACCAC	ACATGGAGAC	TGGCGCAGCC	CTGTTCTGCC
202141	TGGGAAGCCT	TCCCCTGATA	CCCCTAGTTG	GCAGGAGTCT	TCATTTGTTC	TTTTCTAGTC
202201	ACCTGTGCAA	GTTTGTATTG	TTCATGTTTA	TCATCCTTCA	TTCTAGTTGT	CTGTCTCTAT
202261	GTGTGGTCTC	ATTCAGTGGA	CTCTGAACTC	TTATGAAGTC	ATGTCATGGG	TCAGATCTTA
202321	ATAAATTAAT	ATTGTCGGAA	GCTAATGTCA	TGTCTAGAAT	ACAGAAAATT	TATCAAAAAA
202381	AAATATAGTA	TGTTGGCTGG	GCGCAGTGGA	TCAAGCCCGT	AATCCCAGCA	CTTTGGGAGG
202441	CCGAGGCAGG	AGGATCACAT	GAGGTCAGAA	ATTCAAGACC	AGCCTGGCCA	AAATGGTGAA
202501	ACCTCATCTC	TACTAAAAAT	ACAAAAAGTA	GCCAGGCGTG	GTGGTGCCCA	CCTGTAATCC
202561	CAGCTACTCA	GGAGGCTGAA	GCGGGAGGAT	CACTTGAACC	TGGGAGGCAG	AGATTGCAAT
202621	GAGCTGAGAT	CATGCCACTG	CACTCCAGCC	TGGGCGACAG	TGAGACTCCA	ACTCAAAATA
202681	ATAGTAATAA	TAATAATAAT	AATTGTATGG	AATTGAACTG	CTCTGATTGG	AAATAGCTGT
202741	TTTTTAAAAA	ATTATTATTT	TTTAAGTTCC	TGGGTACATG	TACAGGATGT	GCAGGTTTGT
202801				GCTGCACCTA		
202861				AATGTTCTCC		
202921				CTCCCTGTGT		
202981				TTTGGTTTTC		
203041				GGATCTCTGG		
203101				ATTTCATTAA		
203161				ATCATGATTG		
203221				GCAATGCAAG		
203281	TAATGTTTAT	TAAGCGTGTA	CTGTCTTAGT	CTGTTCAGAC	TGCTGTAACA	AAATATCATA
203341	AACTGGGTGA	CTTATAAACA	ACAAAAAATT	TATTTCTTAC	AGTTCTGGAG	GTGGGAAGTC
203401				TGGTGAGGAC		
203461				CTTCCTTGGG		
203521	ATCCTAGTGA	TGAGGTTTCT	GCCCTCATGG	TATAACTACT	GCCCAAAGAC	CCCTCCTTCT
203581				ACATGAGTTT		
203641				GTGCCAAGAA		
203701				CAAAATCCAG		
203761				TGAATCTTAC		
203821				TACTCTTGTT		
203881				TGCTCTGCCT		
203941				GTTCTCCTTT		
204001				TCTCTTTCTT		
204061				TTTTTCTTTC		
			·CIIICIIIC		interrection	.CIIICIIIC

Figure 9 (Page 63 of 74)

204121	TTTTTCTTTC	TGACAGGGTC	TTGCTCTATT	GCCTAGGCTG	GAGTGCAGTG	GTGCAATCTC
204181	AGCTCACTGC	AGCCTTGAAC	TCCAGGGCTC	AAGCAATCCT	CCTGAGTAGC	TGGGACTATA
204241	GGCATGTGCC	ACAACATCAA	GCTAATTTTT	GCATTTTTTT	GTGGAGACGG	GATCTCCCTA
204301	TGTTGCTAAG	GCTGGTCTTG	GATTCCTGGG	CTTATGCGAT	TCTCCTGCCT	CAGCCTCCCA
204361	AAGTCCTGGG	ATTACAGGCA	TGAGCCACTG	CCCCTGGCCA	TTATAACTAT	TTTCATTGGC
204421	TTATCAGGCA	CATGATAACT	ATAATAAATC	AATAACCAGA	ATTTTTAAAT	AAAGAAAGGA
204481	AGGAATTGTT	TCAACTCTTC	CTGCTACCCC	TCTATCCCTC	AAAAGGGTAG	GCTGAATGTT
204541	GTCCTCCAAA	GATATCCATG	TCCTAATCCC	CAGAACCTGT	AAATATATTA	CCTTATATGA
204601	CAAAAGGGAC	TTTACATGTT	TAATAAGTTA	AGAATTTTGA	GATGGGCAGA	TTTTCCTGAA
204661	TTTTGCAGAT	GGGCCCTAGT	GTAATCACAA	GGGTCCTTAT	AAGAGACAGG	CAGAAGAGTC
204721	AGAATAAGAG	AAAAATACTT	CAAGATGTTA	CACTGCTGGC	TTTAAGGTGG	AGGAAAGGCC
204781	AAGAGCCAAA	AAATGCAGTG	GTCACTACAA	GCTGAAAAGA	AAAAGAAATG	GATTTTCCCC
204841	TAAAGCCTCT	GGAGGGGGCA	CAACCTTGCC	AATACCTTGA	TTTTGGCTCA	GTGAAACCCA
204901	TTTTGGACTT	CTGACCTTTA	GAACTGTAAA	TAAATAAATA	ATTTTGTGTT	GTTTCAAGCC
204961	ATCACAGTTG	TGGTAATTTA	CTACAACAGC	AATAAAATAG	AATTAAATAC	AGAGATCTGA
205021	GGAGTTGAGT	AGGATAAGCC	TACTCCAGCA	GGTTATTTCG	GGAGTATGGT	GAGACTCACT
205081	AGGATGGCGG	AACTCAATTA	AGGAAGTCTG	AAGCTGATAA	GCCAGAGAGG	GAAGGCTCTC
205141	ACTTCATTTT	ATAAGGGTTG	CGTCACACTA	GGAAGATCCA	ATAGCAACCA	CAGTCTCAAA
205201	ATTAATGATT	ACAAATAGGA	CACAATTCCA	AGAGTCGGGA	GCCAAGCAGA	AAATGGATTA
205261	GGGAAGACAT	GGATGATATG	AAACAGGAAG	GAGGGGTACA	AGGCAGCTTC	CTGGGAAGTT
205321	GCCAGGGCAG	TCACAGTTCA	CATTCATTAG	GCTGTGGGCA	CCAAATGCAT	ATGGAAAATC
205381	TAGCTGACTT	AACTGAACTC	CTGAAGAGGA	ATGAACACCT	CATTTATTGA	GGAGCTACTA
205441	CCAATTAGAA	TATGTATTTC	ATTTGTTCAA	TAACCCCATG	AGTACAGTAA	CACAATCCTT
205501	GCTTTACTAA	AGCGGAAGCC	AATTCAAAGA	GGTTCAGTGA	CTTGTCCAAG	CTCAGGGAAA
205561	ACACTAGGAA	GTGAATATGG	GTCTGACTCC	ATCACTGATT	TCAGGAGCCC	TGCCCTTTCC
205621	TCCACACCAT	GCCCCCTTGC	TTTCAGAAAA	AAAGGCTTGT	TGACTGAATG	GTTGTATGCA
205681	CAGTTCAAAG	CAGAAACACA	CGATGACATC	TTTTGAGATA	CTCTAACAGT	GAGAACTTGA
205741	AAATGAAGTT	AAAAATTAAG	CGGCAAAACC	AAGCCGAGGC	TTTCTGAGAA	AGTGGGGCCA
205801	AACCTGTTGC	CGTCTGACTG	CCACGTGGCT	CACTATTTAT	CCCTGTAAAA	ATCTGCAAAA
205861	GTATTTGAAA	GGGAAGAAGG	GACAGAAAAC	TCCCTCCTTT	TCCAAGTTAG	CCTTATAGTC
205921	TAGGGCTTAA	AATACTGGTT	TAATGGTGAA	GGTAAGTGCT	TTTCTTCTTT	TTGGGTAGAA
205981	GGATTATTAC	TAACTTACCA	AAGGTCCATT	AAGGGGAGGG	AACAGTTTTA	GGAGAAGTCA
206041	GAGAAAAGAC	ATTAACAGCA	ACATAAGGAT	CTCCATCTGG	TAATATTGCC	TAATTCCAAA
206101	ATGAAGAGAC	TCTCTGAAAA	AGATAACTGA	TTCAATGAAG	ACCCTAGGGC	AAGGCTTGAG
206161	AAGCCACTGG	TACCAATGGA	CACTGTGGAC	AATGGTCATT	TCTCCAAGGA	CGCTGTGAGT
206221	ATTAACTGTG	ATGCTGTGAT	TAGTCAGACT	GGGATTGGCT	GTGGAATGAA	ATACTGATCA
206281	GAACTGACAA	GATTTGTGTT	TGGGACTGTG	GCTAACGAGT	CTTTTCAGAC	TTCTATATGA
206341	ATTTGAAATG	GTCTCTCAGG	AAAAGGAGAA	CATGGCCGGG	CCTGGTGGCT	CACGCCTGTA
206401	ATCCCAGCAC	TTTGGCAGGC	TGAGGCGGGC	AGATCACTTG	AGGTCAGGAG	TTTGAGACCA
206461	GCCTGGCCAA	CATGGTGAAA	CCCTGTCTCC	ACTAAAAATA	CAAAAATTAG	CAGGGCGTAG
206521	CGGCGCGTGC	ACCTATGCGC	ATGCATAGTG	CGCGTGCCAG	CTATTCAGAA	GGCTGAGGCA
206581	GGAGAATTGC	TTGAACCCAG	GATGTAGAGG	TTGCAGTAGT	TGAGATCATA	CCACTGCACT
206641	CCAGCCTAGG	TGACAGAGTA	AGACTCTGTC	TCAAAAAAAT	AATAATAATA	AAAGAAAAGG
206701	AGAACATGAC	CAAAGTTATG	AATAAGACTG	AAGGCAAGAA	AATTGTACGC	TTGTAGAGAT
206761	CACCTAGCTT	GTTGCCCTCA	TTGTACAGCT	AAGAAAAGGC	ACCCAGGGAC	ATTGTGGTCA
206821	GCACCAATTT	CTCAGAAAGA	TAGGCAGATG	ATGAGAGGC	CCTCAGTTTT	TCTAACACTG
206881	AAGGAATTGC	TTCTATGTTT	TCTGGTGAAC	TCCTCCCCAC	TCATCTTGAG	GATTCCAGGC
206941	CAGAAGAATC	CACTTTAAAA	AAGAAACATT	TAAAACCAAT	TTAACAACCA	ATCAAAGGCA
207001	CTTTTATAGA	AATACATTTC	ATTTGCTGTT	GGCCTGTATT	TATGGATCTG	AGAGGGCTAG
207061	ACTGCCAATA	TTGTGACTGT	TTATTATTAT	TGCTGTTGCT	AGTATCTAGA	ATATTATACA
207121				GTCTCATACT		
207181	CCCAGTGAAG	TAACATTATC	CCAATTCTTC	CTATGAAACA	GTGAAAGCCC	TAAGAGTTTT
207241	TGAAACTTTA	CCTGGTTTAC	TCAATTTGGG	AATGGCAGAG	CAGAATTCAG	TCCTTGAATA
207301				CTAGGTGTAA		

Figure 9 (Page 64 of 74)

207361	GGACTCTGGG	CTAACAGAGA	TGAAGCAAGA	CAGGCTGGAT	ATTAGGAGAA	TCTAAGAGCA
207421	ATCTAACGAC	CATTATAATA	AAATCATGAG	TTCTAGACTT	AAAAAAAGGG	AAAAACCTGT
207481	TTTTTTGCTT	ATGCGTATAC	CATAATATTT	ACATTATTTA	TTTTTTTCTC	ANATTCAACC
207541	TATACGGTGT	CAAGTAATTT	TTTTTAATAT	AACATTTTCC	TTTAACTTAA	TTTCNATTCN
207601	TTTTTCTGTG	TCTACTTACA	ACTTTGGCAC	TAGAATTCAC		TACACCTATA
207661	TCTCCTTAAA	GGGAAGGGTT	CTGACACTGT	TACATGTTCT	CAATTITITI	CARAGGIATA
207721	AATAATTATT	CCAGTGTCTC	TAAGTACATA	TCAACCATGC	CAGTGTTCAG	CAAATAGGTT
207781	TTTATTAGCT	TCTGTGCTTA	TTTTGGAAAA	ACATTTCCCA	TTACCATCAA	ACACCTCATAAT
207841	TTAGGATGGT	TTGGTATGTT	AGCCTGATTT	CTGCATTCGT	CTCATCCAAA	AGACCICAGT
207901	AAACGAAGAA	CTGAAATTAC	CTATTGATAC	AAAATCAAAG	TACCATTTCA	DGAAAATAGG
207961	CTTAAGTAGG	GCTTTTCATC	CTTTCTCGTT	AGACAGCAAC	ACACATTICA	AACCATAAAA
208021	AAAGTGATGG	GTTTGTGATA	CAATTCCAGT	AACATAAAGA	CCAACCACAA	AAGAAAAACT
208081	TGTGTTTATG	TTTAATATTC	AAAGCTCAAC	CTAAAAGTAT	TTTTCATTA	GIAGTITIGT
208141	TCTAGAATAA	ATGATTAAAA	CTTGATTTAA	AATATACAAA	TTCTCCTTT	CAAACTTCCT
208201	AAATGGAGCT	ACCCCATTGA	GTTTTAAGCT	TGTGATTAAA	ATATTA	TAATACCTCA
208261	AAGTTGTAAT	AGGTAGAACA	AGCAGTAGTC	TAGGCATTAG	CCCATCTCC	AACAAAGGGG
208321	TGCATCATGT	GGTTTCAGGC	AACTTTTCAA	ATTTTCTACG	GGGATCTGGT	GCTGGCTCTG
208381	ATAAACAGTT	GGGCCAGAGG	ATCTCTGAGT	CTCTTTCAGC	TTTCACTCT	TATCAATAAA
208441	GAGAAGTTGG	TGGGAAAGCT	TTAAGTGGAG	TGTAAGTAAT	TCCACCTCCA	TATAAGATTG
208501	AAGAGTTGCC	TTCAGCCAAG	CCACGGGATC	TTGCATAAAA	ACTCARATCA	TGTACAGTTA
208561	GGTCCAAACT	CTGGGTTTGA	CCACAGATGA	CTTCAGCTAG	CATCTCACTC	AATAGAAAAT
208621	AGCTGAACTC	CTGATATCCA	GATGTTAGCA	AGACTTGGAG	CCCTTCTL	TAGAGCAATG
208681	AACCAGTATC	TGTCCTGGTG	CTCACCTCAT	CTTACTAGCA	ATTECER	GCAGAGCAAC
208741	CATTGTACAA	AACAACAACA	ACAACCIGAI	TAAAATCTCC	ATTGGGCCTC	CATTTGGGTC
208801	TTAGATGGAG	AGATACTATT	CCCACAACAA	TAGAGATATT	TGGAAACACCCAA	AATTCAAAAT
208861	TTGCCATGCT	GATGAAGTCC	AATTATTCCT	CTTTTAAATA	CAMMENAGEAG	AAAACTATAC
208921	TAAAATGAGT	ATCTACTAAT	TATTERCARA	ATCACTTGGT	CATTTAGCTA	CTTCTGAATA
208981	AATGAAGTGA	TCATCCTGTT	TTGTAACCCA	GAAATAGTCA	AAATATAGAA	AGTCACAAAG
209041	CAGTTTCTAT	TCCTGTATGT	GGATGTCCAC	AGCGTATCCT	COTTTOTA	TTGTGTGAAT
209101	AGCATTTTTC	TAATGTAATT	CAATATTCTC	GAAAACATTT	GCTTTGTACA	CTAGAGTACT
209161	TAATCTATCA	AATTGACTTC	CCACACTCTC	ATTATTAGGT	TAAAATAGCT	TCCATCACAA
209221	GCAGTCATGA	GTAATACTAC	AAACCATATT	TTTGGACACA	AMMITTATET	CTAACATTAT
209281	TTTATAATCC	TTCATCCTAA	CCTCACACAT	TATGAATATC	ATTITICATE	TATGCCTTTC
209341	TTTAAATTTT	GTGTGCAAAA	ACAGTCCAAA	GCCTTGAATG	ATARAGIAC	GGACAAGTCT
209401				GGATTCCTGC		
209461	GCAGTGGCAC	GATCTTGGCT	CACTECAACE	TTTGCCTCTT	CCCTTCLLCC	AAGCTGTAGT
209521	CCTCAGCCTC	CTTAGTAGCA	GGGTCTACAG	GCATGTGCCA	CCACACC	AATTATCCTG
209581	ATTTTTAGTA	GAGATGGGGT	TTCACCATCT	TGGCCAGGAT	CLACACCCGG	CIGITITIGT
209641	AAGTGATCCA	CCCACCTCAG	TATCCCAAAC	TGCTGGGATT	ACACCTCTCA	ACCIGACCIC
209701	CCGGCCGATA	CATGTGTTTT	TARRECCARG	GAAATTTCAG	ACAGGIGIGA	GCCACTGCAC
209761				AATTTAGGAA		
209821	TTCTTAGGAT	ATCGATTTT	CCTABAACAA	ACAAATGTAT	TGAATGGAAA	ATTGATGATA
209881	ATTAGTATAC	ממגדדתדמממ	TTARACATCT	CCATATTTAG	CCATCCCCAA	AGATAATTTG
209941	TGTCACAATA	CCTCCATTTA	TTCACAATTC	TAGTAATTAG	AGCCATGAAT	TCTCTTTGCC
210001	CTAGGTGATA	TGAAGACTTT	GTCACAATIG	GCAAGTGTCC	ACATTOTO	TTATAATTTT
210061	GAGAATAAAC	ATTTTAAACT	TTTNAATCTN	ATACATATTA	ACATIGIGIG	TAGCAAACAT
210121	TCATGTTCGA	AGGCACATGG	ANCATTOTTO	TGGTGGTACA	CACCCCACAC	ATGTCATCCT
210181	AGAATGAAAG	GAAAGACCGC	TCTGGAACCT	TCCTCCTTAG	CTCTTCTCT	TACTTTAATT
210241	GTCCTGTCTT	ATGGTCTGCT	ACABGCAATA	CCACTCTTCA	CTCTIGAGCT	CTTCTCTCTCTC
210301	GTTTGATAAA	GTACATGCAA	TOTAL CARIA	AATTCTTCCA	CCTTCGCATG	CARACCACC
210361	TTATCTTTAT	TGAACAGATG	AGGAANTGAN	TGATTAGAGA	ATTTNA NTCH	GAAAGGAGCC
210421	GTCACACAGC	TGGAACTTAC	AGCCACATTT	CCTTTTAACA	ATTIMAATGA	CIAGCICTAG
210481	CCAGTAGTGC	CCCATAAAAT	GTARCTTATA	GAGCTGTGTT	GGGTCNANAC	TTTTACTCAT
210541	GCTAAGAGGA	GGCAACATTA	ÁCAAGGGGAA	ATTATTTGTG	TATTATCTT	TCCATTACTGAL
			MODEEN	VIIVITIGIO	INTIMIGITI	IGGALIAIGI

Figure 9 (Page 65 of 74)

210601	TCTCTCCATA	GATAAAAGAC	TGTCGTAGTA	AAAGAGATTC	AGGGCACAG	GAAACTCCAC
210661	CUCUMAGCQ1	GGIACCAITT	CCCACAGAAG	СТАААТССАС	CCCAACCCCC	00300300
210721	AGGTAAAGCC	ACTGCTCTTG	TTTGCAGGCT	ATGTTAATAA	CCTCA A CCTT	ATTCCGACAC
210781	ATTTACACAT	CTCTGCATCA	CACTGACCCT	TOTIANIA	ACTGGGAGCTT	TAACATTGGA
210841	GCCAGCTCCA	GCCCCTGATC	CTGTTGCTTT	TTCCTTACCC	CCATCALAGIG	TAACATTGGA ATCTGCGAGA
210901	AATTAAGCCA	AATAAGCAAT	AAATCCTGGG	ATCTACCCAC	TCCARGAAATC	ATCTGCGAGA TTTGGGAAAG
210961	TCTTTTTTT	TTTTTTTTT	ACTGAGTCTT	GCTCTCTCTC	ACACCCCCC	TTTGGGAAAG GTGCAGTGGT
211021	GCGATCTCGG	CTCACTGCAA	CCTCTGCCTC	CCGCGTTCNA	CTCATTCTCC	GTGCAGTGGT
211081	TCCCGAGTAG	CTTGGACTAC	AGGCACACAC	CACCATCCCC	ACCTONATION	TGCCTCAGCC
211141	AGTAGAGATG	GAGTTTCGCC	GTGTTAGCCA	GGATGGTCTC	AGC I GAATITI	TIGTATTTT
211201	CACCGGCCTC	GGCCTCCCAA	AGTGCTGGGA	TTACACCCAT	CCCCCACCA	GCCTGGCCCG
211261	GGAAAGTCAT	TTTAAACCAA	CCTATGTATG	AATCCCTACT	ATA ATA TETOT	CACCAAGCGG
211321	CTGGCTCTTT	CTCCTGAGCT	TGGAAACCTC	CAGTAAAATC	CARAMATICI	CACCAAGCGG
211381	CACCACTCTT	ATCTGTGAGC	TTTTTTGGCC	ATTAAAAAIG	ATTTCTTCC	TTTCCCAGAC
211441	ATCTGTGTCT	TCACAGGTTT	TCTCTTTCTT	TCACTTTACT	CCTTTTCTTC	TTATATTTT
211501	GAAAAATCCA	ATCTATCATG	CACATGGGAA	CCCTTTCNAT	ATTCCTTC	AAATAAGCAG
211561	TTTTATGGGG	ATGCTTTTAA	AGAAAAAATT	TGTCCTTTCX	ATTGGTCTGT	GGTTGTTCCA
211621	CACCACATCA	CCTGCAAGCT	TTGTDADADAT	ACTTCTACA	ATATATTGAA	TATCTTCCAG
211681	AGATTGAGTC	TCATTCTGTC	ACCCAGGGTC	CACTACAT	ATTAATTTTT	TTTTTTTTTG
211741	AACCTCTGCC	TCCTGGGTTC	AACTCATTCT	COMOLOGIC	ACATGATCTT	GGCTCATTGC
211801	ACAGGCATGC	ATCACCATCC	CTCCCTAATT	CCTGACTCAG	CCTCCCGAGT	AGCTGGGATT
211861	CATGTTGACC	ATCACCATGC	CANACTOCTO	TTTGTATTTT	TAGTAGAGAT	GGGGTTTCAC
211921	CAAAATGCTG	AGGCTGGTCT	CETCACCCAC	ACCTCAAGTG	ATCCACCTGC	CTTAGCCTCC
211981	AGTTGAACAT	GGACTACAGG	ACCACCTACE	TGCACCCCAC	GTAGTTTTTT	TTTTTTTTTA
212041	ATTACACTAG	ATGTGAAGGC	AGGACCTAGT	GACACATAGC	AATAACATTT	CCAAGTAGAC
212101	AGAGAATCCT	GGAATTAGTC	AAAGIGCICA	TTTAAAGTAC	CATCTCTCAA	ATGTATTAAA
212161	AAGCTTTGTG	TGGATGTGCA	CTCCATAAC	TCAAAGGCAG	CTCGTTATGT	ATAAACTCTC
212221	TTTTATTGAC	ATAAACAAAT	TTATOTOLOG	GATGGGACTA	TTGACTTACA	GCCCAGGGAA
212281	ATTTTGGAGC	GCTGAGAAGG	AAATGGGGT	GGCTCTGCCA	CTGTCATCCC	CATTCACTTC
212341	TCCTATCCCC	AATATGACAT	CCCLEA	CATGTGGGTT	TTCTCTATTT	ATCATGTGTT
212401	TGTCTTGAAG	TTGAAAGATG	CCATATTIG	CTTTACTTGG	TTATAAGATC	CCATATTCGC
212461	AAAAAAAAAA	CCAACCAAAT	CATCTCTCACAA	AGTGGGTTTG	TAGTGCTGGC	TATTTTGGTG
212521	AAAGGAAAAG	AATGAGACTT	CAIGIGICAT	CCAAAGTTCT	ATCAGATCGA	GCTGTGAGAG
212581	GGTCCAGATT	AAAGGGGTCT	P.C.C.T.T.	GCTCACTGCA	TACATCTGTG	TTGTTGTCTA
212641	ACCATGATAA	TCTGTTCATT	ACGCTATGGG	CTGGCTCTTA	TCATGCACTT	CTCAAACTTC
212701	CAGCAGCAAG	CGCAGCGTGT	TOCOTOON	ATTGCGATCA	TCGCCATGGT	GAACACCACT
212761	TCCAGCATAT	GTCTATCTAA	ATTTCATAGA	GAGGGGCCTG	TTGCAGATGC	CTTCAATAAC
212821	TGTTGAGAGA	CCATCAAGGA	ATTIGATACA	AAGGTAAGTA	TGATGGAAAA	TAGGGCTCTT
212881	ATACATTTCC	AAAAACTTTG	AAAGGAAGGC	ATAGATCTTG	ATTCTGTGGA	GIATGGAAGT
212941	CTTCDATAAT	AATGACAAAT	TAAAACTGAC	TGGAACTATT	TTTCTTTGAG	ACATTGCTTA
213001	AGAGTTAAT	AAAAATAAGA	TARRATTGAG	GTTATTATGA	TTATAAGGTG	GGGGAACTGT
213061	TAGGTATTAC	GTGAAAAATT	TAAAAATGGA	ACAGTTTATG	TGATGTCTTC	AATGAAAAAC
213121	TTATTGTTTC	CTGGGCACAT	TATATAGGT	TACTCAATCC	TATTCAGTTC	TCTGCCTGTT
213181	CGATTCTTCT	TGAGCAATTT	COLLEGE	TAAATTCTAT	ATAACCAATA	GAAATGCAAA
213241	TOTOCACTOA	CCATAGCTTT	GCAAATAAAT	TTTGCCAAGA	GAAAAATCAG	TTAAAACTTT
213301	TGAGATACAG	CCTCCCAGTT	GAATTAGCCA	ATTTTGCTGT	TTGTTTGTTT	GTTTGTTTTT
213361	GCAGCCTCCC	TCTTCCTCTG	TCATTCAGGC	TGGAGTGCAG	TGGCATGATC	TCAGCTCACT
213421	GEAGCCICCG	CCTCCCGGGT	TCAAGAGATT	TTCCTGTCTC	AGCCTCCCAA	GTAGCTGGGA
213481	ACTAGGGGGGG	ATGCCACCGC	GGCTGGCTAA	TITITGTATT	TTTAGTAGAG	ACAGGGTTTC
213541	TTCCCATTA	TCTCGAACTC	CIGACCTCAG	GTGATCCACC	CGCCTCGGCC	TCCCAAAGTG
213601	CONTROCTE	AGGTGTGAGC	CACTGTGCCA	GGCTCTGCTG '	TATATTTAAA	GTCTATTTCA
213661	CATIGUIT	CTGCTTGTGT	TATGCGTGAT	TCTTTGAGTT	TTCCTTTGAA	CCAGTTATAA
213001	TTTACATOR	ACTTCCTCCA	I TAATCAATG	AGITAAATAA	AATCTTTGTT	GTATGTTTAT
213721	TTCTDCTCT	TATGAAAACC	ATGAATTTAC	CCAATTAAAA .	AAATTATCCT	TTAAATTATC
	TIGINCIGIA	CATTTCCÇAT	GICATCCCTA	TAATTCATGA '	TTAATGATTT '	TATTACATTG
					•	

Figure 9 (Page 66 of 74)

213841					CCAGTCTTTC	
213901	CCGTCTACAT	ATCCACACTG	AGTAGATTCA	CTACTCAGGA	ATCTTGGACA	CCTTCAAGTT
213961	GCCAAACATG	CAGTGTTCAC	TGGACATGCT	GTGTTCCTTC	AGAATTTGGG	CCTGCTTCTC
214021	AGCACACTCA	CATCTGCTAT	CAATGACCCA	TGGAAAGTTT	TTGCCCTGAG	CAAGCCAGAG
214081	TCCCTGTTAG	TTTCTTCCAA	ATGCTACAAG	TTCACTTTTG	CTATTTTTC	CGATGAGATA
214141	AAATTTTCCT	TTTTGACTTT	CTACAAATCA	TAGTCATTTT	TCAAGGGATA	GTTCAAGTAT
214201	TGCTTCCTTT	CTGGGACCTT	CCCAAATTAT	TATTTTCTCC	TCTCAAAGTC	TCTGTTTTAT
214261	TTATGTTCAT	CCTCAAATCT	TGATTCTCAC	ATGAATCATA	TACCTTGTAT	TATTTATAGT
214321	TTTTTTGAGT	AGGTAAAATA	TTTCATATTT	TATATTCTTT	GGCTCTCTAC	TTTATAGCAT
214381	GATGCCAGAT	ATTTAGGGGC	CTTACTGCAT	TTATTTTTTA	TTTTATTTTA	AAATCTATTT
214441	TATTTTTTAT	TTATTTATTT	TAAAATCTAT	TTATTTTTAG	GTAAATATTC	AGGTAATATA
214501	ATTTATGTAA	TTATTTAGGA	ATTTTAGGTA	GTTATTTTAA	AATAATTCAA	ATTATTTATT
214561	GAGTTATATC	AGAAGAATGT	GATCTTATTC	ATTTGTAATA	TGTGTTTTAG	GAACTCAGTT
214621	CAGCCAGGGC	AGACCATAAT	TCCCAAACTT	GACTTTTCTT	TTTAATTAGG	CACTGATTTT
214681	GGTTAAGAGT	TCAGTAAAGT	TTTGTGTGTG	TGTTTTAAAA	AATTCTTTGA	TATAAGAGTC
214741	AAGATGTTAC	TCAACTTTTA	CTAGAAGCAA	AATAGAGGAA	GTGCTTTCAC	AGATGAAATA
214801					ATCTATATAA	
214861					AGACAAGCAA	
214921					ACAAAAATCA	
214981					AGATTTTCAT	
215041					GTATCAATGG	
215101					ACTGACTCTG	
215161					TGCTGGTTTG	
215221					AGTGATTTTG	
215281					TCTCATTCTT	
215341					TTTCATTTCA	
215401					CACTTGAACG	
215461					ATAGCTTTGT	
215521					GGAAGAGAGA	
215581					TAGGAAAGAT	
215641				•	GAGCTAGGTT	
215701					CCTTGTCGGA	
215761					TATATATACA	
215821					GTGGTAAGGA	
215881					TACTATTCTA	
215941					AAATTCTTAT	
216001					TACAAATAGC	
216061					GCAGACATTC	
216121					CTCTACAGCT	
216181					GAGCAGAACT	
216241					GCAGACTTTT	
216301					CATCTCCTCT	
216361					ACCTTCTTCC	
216421					CCCGCTTGAG	
216481					TTGCATGAAT	
216541					GTGCAAGCAT	
216601					ACTTCCCTCT	
216661					CTTGTGCCAA	
216721					TTTTTTTTT	
216781					CAGGCCGGAC	
216841					CCATTTTCCT	
216901					GCTAATTTTT	
216961					GACCTCGTGA	
217021						AAAACTTCCT

Figure 9 (Page 67 of 74)

217081	AAATCTTATA	ATTATTATCA	ATTTATCCTC	AGATATACTT	CCACGTACAT	TGTAGTTTTA
217141	TTATATTTAT	ATTTTACATO	TTTTTTTCA	AATTGCAGTT	TGGGACCCAT	TAGTGAGTCA
217201	TAAAATCCAT	TGAGCGGGTT	AAAATCATTA	ΤΤΤΤΟΔΑΔΑΤ	TGAGTAGAAT	AGAATAGAAA
217261	TTGTTGGAGT	GCATTGGACA	TGGTAAAGTT	AAATATCGAT	TCATCAAACC	AGAATAGAAA
217321	GCATATGTGT	GTGGTTGTAT	GTACAAGTGT	TTATGCATAT	TEGTETETET	ATCGTTTGAG
217381	CCTGTAAAAT	GCATTTCTTA	CTATAGGTCT	CTCTCDAATA	TCTCTCTTCT	TGTTTTTTAA
217441	TGTAGACTTC	CAAAGCCTAC	ATGGCATTTC	ACTAGTGACA	TGIGICTIGT	TGTTTTTTAA
217501	TCTCTCCAAT	TGGACCAGAA	GCTCTTTGAG	GGCAGGGGCT	CTATOTTA	TTCACATTTT
217561	AGTCTTTCAT	TTCCTGCCCC	TAGCCTCATA	TTAGATCATG	GIAICTIACC	GATTTTTGTA
217621	CAAGAAAATG	CTAATGGGCT	GTGATAGCAG	AGAGTTACTG	TCACAAIGCA	ACTGTAATCA
217681	ATTTGGTCAC	ATTGGTGTTG	AGGAGCCATT	GAAGAATCAG	1GACAAACTA	AGGGATTTAG
217741	GTTAATTTTA	ATTATATCAT	ATTACTTTAC	TGGGGAAAAT	AGAGIGIGIT	ACTATTATTT
217801	AAATACTCTC	ATTGCCCAAT	AATTCTAACT	CTGCCACCTC	CIGIGAGCIA	TTTTAGAAAT
217861	GGAGGCCACG	AAGTCTCAGC	CTTTCIAAGI	TTCATAAGTG	ACTGTTGGGA	CATTGTTTAG
217921	AGGGTCAGCA	TTTGGATCCT	TCATCATCCT	CTGTGTGGGG	TITTTCTCCC	TTTTTCCTTT
217981	GAGCTGGCCT	TTTATCTTCT	ACATCATCCT	TGAGTCACTT	GGACTAATCT	CACAGGCCTT
218041	CCATTTCCTG	AGCATCCATT	TTGGCACCTA	CACCACCCAC	TCTCTTAAAT	CCTAATGCCT
218101	AATGTCCTTT	ATCAAATGGA	AGATGATAAA	AAATGTCAAC	ATTOTTCCTA	TATGAAAGAA
218161	TAGTCACACA	ACCTGATTAA	CACCTTCCTC	GTGGTTCTGG	GGTTGGTATC	ATTTTTAATC
218221	GAGGAGTTGA	CTATTCACAT	GGCACCCACC	GIGGIICIGG	GAAGCCACAC	GCAAAAGGTA
218281	CAAGCACCTT	CTGCAGAATC	TCTACCACCA	GACTTGTGAT	GCAGTCTTGT	CCTTCCATAT
218341	ATGTCAAAGA	TAGTGAAGTC	CATTOTACCACCA	CATCTGAAGT	GCCTGCTATA	TGCAGTTAAG
218401	TOTOTOGA	ATCCCTTTCA	CATTTTCAAT	GTGTCTTCAT	ATTTCATTAT	AATTATTATT
218461	ATGTTCCCTT	CCCCATCCCC	CCTGTTCTCT	ACCAAGTTAA	TCTTGCAAAG	TTCAATTCAA
218521	TTTATCATAT	TTCCTCTCTA	CCTTCCAGGG	CTTACCCTGT	CAGATTCTGG	CATTCTCTCC
218581	CCACTAGACT	CTCARACCT	GGTTATGTTG	GTGTGTAATT	ATTTATTTCT	CCTTTTCTTT
218641	TCATCATGGT	CCCTCATOTT	TGAGGCAAGG	AATCCATTCT	ATGTTTTCAT	CACTTGGGTG
218701	GGGGATTTAA	DCDDDDCTDC	TAGCTTTAAA	ATAAAAGAAT	CAGTGAATCC	AGTAATTAGA
218761	AATTCCAATA	AGAAAACIAG	TTTTCTLAGAAT	CTTTTAACAT	AGAATGTTCT	TCAAATAAGG
218821	TTAAATATAG	TCCTGGCCTC	AATCCCTTTTC	TGATTTTGTT	TITATAGCCA	AATGGTGTCA
218881	ACATGTTAAC	CACCTATTCT	ACARAGETTIC	TCATTAATGA	TGCTAATTAT	TTTGGTTTGT
218941	CTTCAATACA	AATAATACTC	#CAAAAATAT	TTCTTTTGGG	AATCCATAAT	GGATGTATGG
219001	TTTCATCGAA	CCTTCTTTCC	TCTCTTGTAA	GTGCATTGGA	AATTTTTCCC	TGCCACATGA
219061	AACAACACAA	CTTATCTCTC	IGTATGTATG	ACTGCAAACC	TGACTATTCA	GATCTTCCGC
219121	ATTICANCACAA	TTARACTAR	CATTAAGAAG	TTGCTGCCTA	AAATACATAA	CACTGTAATC
219181	GGCTCTGACA	TTCACAAAT	TAATCAGCTA	TGCAATGCCA	CGCTCCTGTT	ATCTCCAGAG
219241	CACCTTTCTA	CAACCAMMOA	GIGGCTTCT	ATTTGAGACG	TAATATCTAA	AAAGCTTTAA
219301	GCATTAATTC	ATTACTOTOR	AAGAAAGAAT	GGGAACATTT	AGGTCCTTAT	GGTAGAATAA
219361	CCCAGTAAAC	ATTAGIGIGI	AGAAGGGAGA	GGCATGCCAC	TTCAGAGGAA	ACTTCCTTCC
219421	GTGTCTGCTG	TOTOCTATO	TTCACACTAAT	TTTATCCCTT	CTTCCCAGGT	AGCACTGGCT
219481	TARGTGTTAG	CCAAAACCAC	CLCACAGTGA	TTTATGATGA	CCCCATGCAT	CACCCGTGCA
219541	CTTCTACCTC	TCCCCCATCC	CACATCCTGT	CCTCACTGGC	TCAACAGGTA	CAGTGCACAC
219601	ACCATCTTCC	CTCCTCTAAT	AGAGGTCTCT	AGGGCAGGGT	GTGGATCTCC	TCTGAGAGGC
219661	TGGACCACCT	CTCCCCAMA	ACTCATGCTG	ATTAGATCTT	TCTTTTCAGC	CCAGTTCTCC
219721	GGGTTTTTTC	ACCCATAA	AGGCGATGGT	CACATGCCTA	CCACTTTGGG	CCATTTTCCT
219781	CAGTACTCTC	AGCCAT LICT	GGTTATGCAC	CATCATCCTA	ACATACCTAC	CAACGTATAT
219841	DTCDTDDTCC	CICCAIGITA	ACATCAGAGA	TGTGAGTTTA	CTTCCTATAC	TTCTACGAAA
219901	TABCCATTA	TARTAAGGAG	AAACAGTTCT	GTGTTACCTA	TTACATTCTG	GCTTTACATA
219961	ACACATIAA	111AACCTTC	ACAATGACCT	TGAGAGAGGC	ATTGTTATAA	TTCCCTTTTC
220021	ACAGATGTGG	CTCCTCCTCC	CTTAGAGGTG	AGATAACTTG	CCCCAGGTTG	CACAATACTA
220021	AGTGATAGAG	CIGCIGCAGC	ATCCATATTC	TTAACCACTA	TGCTATACTA	CCACACCAGC
220141	TAATTCCAGC	ACTICITITA A	GAAATAATAT	TGCTGGGCCA	GGCATGGTGG	CTCATGCCTG
220201	TAATTCCAGC	TATCCTTTA C	GUCGAGGCAG	GCAGATCATG	AGGTCAGGAA	TGCAAGACCA
220261	GCCTGACCAA	TWIGGILIAC	CARACTATCAT	CTACTAAAAA '	TACAAAAATT	AGCCAGGTGT
	GGTGGCAGGC	VCCTGTWWTC	CCAGCTATTC	AGGAGGCTGA (GACAGGAGAA '	TCGCTTGAAC

Figure 9 (Page 68 of 74)

220321	CCAGGAGGT	G GAGGTTGCA	T TGAGCCAAG	TCATGCCACT	CCNCTCCNC	
220381		* CCGITICMM	M AACAAAAAA	ነ ሮሮሽሽሮሽእአክምቱ	*	
220441		u rackaciic	i GGCCCTCTT) TCTCカCカペカペ	TOWNSON-	
220501	GGATGCTAA	T TTTCCCCCA	A ACAACCCAC	GTATCATGGG	GITCTTTA	GTGTGAAAAA
220561		u cumutitic	3 IGCTAACCT	, TOTOTANA	***********	
220621	CCTTCAGAG	T GGAGTTCTG	CCTCCCTGC	TTTTATTGCT	TACTCTGTAT	AAACTTCCTT
220681	AGGAGGTCA	G CTGGCAGAT	TCCTTTTGTC	CAGGAATCTT	GCTGCAAGCT	GTACAATTTT
220741	AAAGCTCTT'	T TCATCTCTTC	GTAAGGATA	GCGTGTGGGC	CTCAGATTGA	TCACTGTGCG
220801	CTGCACATG	G TCTCAGAGGG	TTCCCTGACA	GCATGTCCTC	CCATTTAACC	AATCCCTTTT
220861	CCATCAATA	T GTGCTGTGG	CCTGCCCTTT	GTGGCCTCCA	ATTGCCCAGG	GCTCCTCCTT
220921	TTGCTGATA	TTATTCCTG	GACCAGTAAC	CTATGTGACT	GITACGTGAT	AACCATTATT
220981	TTAGATATCO	CCCCCAGGT	ACACCTCTAC	CTGTTTTTC	CAGGGTTTAT	CATCAACACC
221041	GAGGTGTTA	ACCTCAGTGG	TCCCCCTCAA	CIGITITITE	CCCTCCTCCA	GACCCCTCCA
221101	GGCAGAATG	CAAATAACTA	CANATATOTO	ACTCTTTAAT	GTTACTGACA	TTGCACTAAT
221161	GCATTTTTAC	AACAACAATT	TCCAATAICIG	TCTGTGGCCA	TTTTTAGAAC	AACAAATGTG
221221	AAGCTTCCCT	AACAGAGATT	CAACTCTCTA	GCCAGTAATC	ATTTTGACAA	AAACCTTCCC
221281	TGGAAAAGTT	TOCATOGRATI	GAACIGIGIA	TGCTGGGAAA	AGGCCCACAC	ACAGGTGATT
221341	TATATATATA	TATATATATA	TOTICATATI	AGCTACCACA	TATATATATA	TATATATATA
221401	AGACTTGCC	· ININIMIAINIA	CATCELLE	TACAGTCACA	ATAAGCCAGC	TCCTGTGCCA
221461	ATCCCCATTT	TATAICAACA	CATCTAATCC	TCACAGTTAT	ATTAGGTAGG	CCCTATTGTT
221521	CACATAAAGG	CACACCCACC	AAGGCTGAGG	CACAAGGAGG	TTAAATGGTG	TGACTATGGT
221581	CCCGTTGCAC	. VVVCLCCUCA	ATTTGGACTG	GGGGAGTCTG	GCTTTGGAGT	CTGTGTCCTG
221641	GAGACTGCAT	TECTECETE	CTACACTGAG	CAGCCAGGGT	AAAGAAACGT	GGTTCCCAGA
221701	or to to to CAL	100100100	LIATIGACTT	GGTAGATTGG	TAATTTCACC	
221761	CATTAGAGGT	CIGAAIGICI	TTAGGTGAAT	GAAAAACTGC	ATTAAGCAAA	ATGACTTTGC
221821	TAAAATCATT	GAALIGCALL	AAAGTTGAGT	TGCTGCAGAA	GCTGTAGGTG	GCTTTCTATA
221881	GGGGATTTGG	CCTCATCCCA	CCTTCCCATA	GATATGCAAG	TTTCCTCATG	GGAATCTCAA
221941	AGGTTGGGTC	ACTETATEGEA	GGAATCATCT	CTTCCACTGC	CACTGGATTC	CTCATCAGTC
222001	ACACACGGTG	CTCTAAACAM	ACATCTTCAA	GTGGCAGGTA	TTGTTTTAGG	TGTTGGAGAT
222061	AATCAGACTC	TCCTACCTC	CIGGATGGCA	ACACAATTAC	TCTATTTACA	TGAGCCTCTA
222121	GATGAATAGA	TCTTACATTC	GATTTCCCAG	AGGAAGAAA	ATATAAGCTT	ATTTTCTCAA
222181	ACTTCCTACA	CCTACATTG	ATTAAAATGA	GCTGTTCCGG	TGCAGAAGAC	AGCACGTATG
222241		GGIACAIGAG	CATGAAACAG	TTCTTAGTTA	TCACCACAAA	
222301	TO T CHARGOAM	INGCAAGAGA	CGAAGACAGA	GGGGCCAAAAG	AACATCATCA	
222361	create that c	CAATITIAA	AAAATCACAA	AAGGGAAACA	ARCTOTOOTA	~~~~
222421	INCOMINATIO	MAIGICIGGA	AACAGATCGG	CTGTGAGACA	TTCCNACCAC	
222481	TOTITOGAMA	IGCAGGCICA	TGAGGAAGAT	GAAAAGACAG	ACCCACCOAC	
222541	HOMETONE	ACCAMCTIAC	AAAGAGAAGT	V-L-M-L-L-D-L-L-L	CTACAMMON	
222601	arreceyod!	INMINITION	CTAAACTGCT	AGGAATCCAC	アクサクカ クマカマカ	\
222661	TOTICITACIA	GGGCIIICIG	AGGAGGGTCA	CACAGAAGAG	~~~~~~~~~	mas =======
222721	TECHONIGG	LIMINGIGAT	AGTTGTCAAC	ACCCAATACA .	~~~~~~~	
222781		THE PARTIES	ACAAAAAAA	AAAACACACA	7C1C1C111C	
222841		INGOCKINAL	TITAAATGAG	י אידיאידיאידאידאיי		~
222901	OLD DOGGETT T	MOIGIALLI	ATTTTTTTTT	ΑΔΑΓΙΑΝΤΑΝ .	CCATCTCARG	
222961		IGGCCWIIII	GITTCCAATA	ע ממיד מיייידיים (מייידיים או	TO A STATE OF THE PARTY OF THE	***
223021		TIMINITUCE	IGIGATCAAC	ATTGCAATAC :	1 C 1 1 C T C C C C 1	
223081		GWWGGWWC I I	GIGAGATTGA	TCATTTTCTC		
223141		LIGGAGGAAI	GICTITITCC	TGTCTGCTGC :	CTCDACATO .	
223201	*CITITACCI	CACGIIIGGA	CAAGCAGAAC	TTCAAGACTG (ここころ カカスカー	CCACCCC
223261		AGGACATAAA	GTTACAAACT	TAAATGTGGT	CTCRCCRTC .	
223321	"CHITITIA	CITCICICCA	TATTCCTGAC	ሮልፕልርልሮፕሮል ረ	CACTTOTAL .	
223321	TOTOTIAGIC	116661666	AGCCTTTATA	ልርልሮልሮፕሮልቱ <i>፣</i>	CTTCCCACC	
223441	ATTCIGANIG	WALIGGICIG	GGGTGGAACC -	ר אכשרארידאר יו	. Deminstration (4.1)	M >
223501	CAGGIIICIA	GCMIGCGCC	GGGGTTGACA	ACAGCTGGAC 2	A A CTTCS B B .	
	TGTGGCCTTT	GAATTTTCCT	CATTGGAAAG	TACTAAATAA A	TAAAAATTC /	TGTGAAAAT

Figure 9 (Page 69 of 74)

223561	Chmch cmah m		,			
223621	GAICACIGAT	AAATATCTTC	ATGGTGGGGC	AGGTTATTGG	ATGCAGAGAA	GATCTGCTCG
	GAATTGTAGC	CATATGTTAC	AGATCTCAGC	ACCGATCAGA	ACTGTAAAGC	TATAATCCCC
223681	AGAATTAAAG	TTTTTATTAT	TTTTTATACA	TTGTAAAACA	TAGACGTTTA	TTTATGTGAT
223741	TAAATTCTAT	TAAAATTTAC	ATGCTAAAAT	AAAATAGACC	ATTTTCAAAT	TATTTAGATC
223801	CAGATATTTC	CATCAGATTA	AACAGATATT	TATTTATCCT	AGCCCAATTG	CAAGAGATTA
223861	ATGATGAGAA	AATGACCAAT	ACAAGATTAA	ATAAATGAGG	TTAACTTAGA	AATCAAGGAC
223921	AGAGAAGATA	GAACTGGAAA	GCTTGTATTG	TGAGAAGAAT	GAATGTGAAG	GAAGGCAATG
223981	TAGACACTTC	CAGAAGGGAT	AGCAATATAG	TTTAGACCAT	ATAATGAAAA	TTGGAGAGAG
224041	ATGACAGAGA	CACTTTCAAG	TGAAATGACA	ATTTATATGG	GGGAGAAAA	TATTGAAGAC
224101	ATAACAAGAT	GAGAAAAGGC	ATAGAAATGT	ATCACATACA	AGGCATAGAA	GTGTATCACA
224161	TACAAGAGAA	GTTCCTTTTG	AGCGTAGAAA	AAGATAATTT	AACCTTCTTC	ATATTTTTCT
224221	TACTTTCCCA	AGATACTCAG	ATAGGCAGCG	TCAACTCTAA	CAGGAATTAA	TTTGGCTCCT
224281	AACACTTAAG	ACATATCCTT	TAGTTTGTCT	CCTCACACAG	AACTGATTCT	GGTTTTGCCA
224341	CAACATGTCT	AGAGAAGAAG	TTCCCACCAT	ATTTTAAATC	CTATTAAAAA	ACTGCTTGGA
224401	CAAGAACCTT	GGGCTAATTC	AGCAGATGAA	GAGAATCTCC	TAATGCAAAT	CAATGGGTAT
224461	TTTTGAGCAA	GTTTTTCAGA	AAAACAGAGT	GTCAGGCCCT	GAGGGTGGTA	CTAAGATGAG
224521	AACATTGATT	TTGCCTTCAT	GATATTGACA	ACACAAAGAG	GAAAGGGGGT	TTGCAGAAAA
224581	CTAAAAGAAG	AAGTAGAAGA	AAAAAGAAAG	ACATAGTATA	ATAGGTAGTC	AAATTATGTA
224641	CAGAAAAAAG	AGGAAAAAA	ACCAAAAAAG	GGTGGGGGAC	AGACAACCCA	ACTAAAAAAT
224701	GGGCCAATGA	CTTGAACAGG	GACTTCATAA	AAGAGAAAAT	GTAAGTGGCT	CCTTAACATA
224761	TAAAAAGATG	TTCAACTTCA	TTAGTCATTA	CAGAAATGAA	AATCAAAACT	ACAATGAAAT
224821	ACCACTATAA	AATTAACTAA	TGGATAAAAT	GAAAGGAGAT	GGAAAACAAA	ATGTTGCCAG
224881	ACATGTGGAG	CAACTGGAAC	TTTCATACGT	TACGAATGTG	AACTTTGGAA	AGCTGCTCGG
224941	CAATATCTCC	TAAAGCTAAA	TGTACAATTC	CAGTGACTCA	GACATTTTAC	TTAGAAATGC
225001	ACATATACAT	CCATAAAACA	TGTACAACAA	TGTTCATAGG	AGCACTATCT	GTAATAGCCT
225061	GAACAGGAAG	TTGTCTGTTA	AAAAAAGAAT	GAGTAAATAA	ACCACGGTCT	ATTTGTATAG
225121	CAATGAGAAT	TAACAGACCC	CAATATATAA	TAGATGAATG	GGTCTCATAA	GCACAATATT
225181	GATTAAAGGA	AGACAAAACG	CACATTCTTT	TAAAGGTTTA	TAAAATACTT	TTTAAAAACA
225241	GCTACAACCA	ATCCGTCCTG	TTAAAAATCA	GTGAGCGATT	TCCCTTGTGC	AGGGATGGGG
225301	GTTGTGGCTG	GATGGATGGT	ACTTAAGAAG	TGCTCCTGGG	GTACTAGAAA	TATTTTATTT
225361	CTTGACTTGG	ATGTGTGTTT	ACTTTGTGAA	TATTGTACAT	TTATGATTTG	TGCACGTTTA
225421	TGAATGTAGA	AAATAAAACA	GAAAGCAAAT	TCAAAGTATC	ATCCTTTTGA	GAGCTTCTGC
225481	TCTGACTTCG	TTTTGACCAA	TGGAGCAGTT	GGGAAGGGGT	CTTGGTCCTT	CGGTCCTTTG
225541	CTTTTTTTT	TTTTTTTTTT	TTTTAGACAG	AGTCTCACTC	TGTCGCCCGG	GCTGGAGTGC
225601	AGTGGCTCGA	TCTTAGCTCA	CTGAAAGCTT	TGCCTCCCGG	GTTCATGCCA	TTCTCCTGCC
225661	TCAGCCTCCC	CAGTAGCTGG	GACTACAGGC	ACCTGCCACC	ATGCCCGCCT	AATTTTTTGT
225721	ATTTTTTAGT	AGAGACGGGG	TTTCACCATG	TTAGCCAGGA	TGGTCTCGAT	CTCCTGACCT
225781	CGTGATCCGC	CCACCTGAGC	CTCCCAAAGT	GCTGGGATTA	CAGGTGTGAG	CCACCGCGCC
225841	CGGCCCCTGG	TCCTCTGCTT	TCATGTTCTT	CTTGGTCCTG	TTCCTCCTCC	TCTTTTGTTG
225901	GAACTTCCAG	TATCAGAGCA	GGAAGGAAGG	CAATGGGTCA	ATCGATGCTG	TCAGCTTTTG
225961	GATCAAACTG	CAAGTTCTCA	AACAGCAAAA	TTAATGAGCT	CAGGCTTTGA	AGAAACCATG
226021	ACCCTGAAAG	CATCAGTTGC	TTCCAATTGC	ATCAGTTGCC	ACGGGTGATA	AGAACAATGA
226081	TGACTCAGAA	TGCCTAGGTT	TTCCCAGCAG	CTTCTCTGAG	GTTTTCCCAG	CAGCTTCTCT
226141	GATTGATTCC	TGACAGATGA	CTTCGGTGTG	TCAGACTTTC	AGGGTATCTT	TCCTTATGTG
226201	ATGGTTTGAG	GAAGAGTTAC	CATTCACATT	CCTAATGGCT	TCAGAATAGA	TGCAATTGTG
226261	AACTGATAGG	AAACATTTCT	AATTCATCTC	CCCTCCCCAT	CCCTAAAGGA	TTGTTTCTAA
226321				CAAATAGTTT		
226381	GAGATGACTT	ACTTTTTCTC	CTTGACTGTT	AAATATTATG	AATTATATTA	ATGTATTTCT
226441				TGATGTACGA		
226501				TATATGAGAT		
226561				TGGAGCTTAT		
226621				GAAAAAGATT		
226681				CATTGGTTTG		
226741				TCAATTTCCT		

Figure 9 (Page 70 of 74)

226801	GGGTTTTAT	A TTTTTCTTTC	ATCAATTTT	G ACCATTTATO	: TTNTCTTCCN	CCATCATON
226861	* * * TACACAC	- IAIIIAAAG	LATATTTGCA	A AAATTCAACI	CTTTTNTCNC	CCTATCTTTT
226921	TAATAATAT	A TTCATTTAT	CTATATCTG	A GGTTTTAGCT	TOTTTALCAG	TOTALCITI
226981	TTGCATGTG	r GCTTTCTTTC	TCCTTCATT	A GACTACTTAC	TCTTTGIACT	TOTGACCCAA
227041	TAGCTTGTC	TTTATTTATT	TACTTATTT	TTTTTCACAC	CCACTCTCA	ATTTTAAGAA
227101	AGGCTGGAGT	GCAGTGGCGC	GATCTCGGC	T CACTCCAACC	. GGAGICICAC	TCTGTCACCC
227161	GATTCTCCTC	CCTCAGACTC	CCGAGTAGC	CACIGCAACC	TCCGCCTCCC	GGGTTCAAGT
227221	CTAATTTCT	TATTTTTAAT	AGAGATGGG	TTTTCCCATC	TTCCCCARCCA	CCATGTCTGG
227281	CTCCTGACCT	TAGATGATCT	ACCCACCTTO	GCCTCCCAAA	CTCCTCCCA	TGGTCTCAAA
227341	AGCCACTGC	CCCAGCCCTG	CTTGTCTTT	TATTTATAT	TTCATTACCT	TACAGGCATG
227401	TCAAGCTTAT	GTCCTATTTC	CCTTTGCTT	C DOTTONTATA	NATITATION OF THE PROPERTY OF	TTATCTTTA
227461	ATTTATTTT	CATTTAATTA	TGAAACAGG	TABACCTTAC	ACCARART	TGGATAGTTT
227521	TCCACTTTTC	TGGGCAGATT	ACATTTTCC1	CTCTTCTCCT	AGGAAAATTG	CTCCTCTAAG
227581	AATGCTTTAT	TTCTCAAGTT	' AATAACCTAT	TATACTARARA	CCCAAATTCA	TIGITCITIT
227641	TTTTTTTTT	TTTTTTTTT	ייייייייייייייייייייייייייייייייייייי	CATACACCA	AGTGGCTGTT	GACTCTCAGC
227701	TGGTCTGAAA	CTCCTGGCTT	CAAGGGATCC	TCCTCCCTTC	CTTGCTGTG	TTGCTCAGGC
227761	GACAGACATO	AGACACCATG	CCCAGCCATC	TOTOTOTO	GTCTCACAAA	ATGCTGGGAT
227821	ACACACTGAG	GCATCCTATC	ATCTCACTACT	TOOTTO	TATATATAAT	AAGAAAACAG
227881	CTCTGACCTT	TTGCAGTTAA	TGTATTAATT	TGGIIICACT	ACTGTTCTCT	GGAAGTTTTG
227941	TAGCATTTGC	ATTCTGTTGG	GTATTATACT	TIGCATIGAG	TAGTTTCCAT	AGAAGAATTA
228001	CTGAAACCAA	GATGAGGCAA	GTGAGGTCCC	TITCACTGTT	ATTTGAACAT	AATTTGAGGG
228061	CTTAGGCTCA	TGCAAGAACA	GAATTCCCAC	CAGGAAGCAA	TATTTAAGGA	GGCATCCTTT
228121	GCTGGACACT	TGCAAGAACA	TTACCATACC	AIGAGAGTGA	GTGCCTCCTT	AATTTTGAGT
228181	TTTTCATGTC	TCTTGCTCAC	COTTOTTO	CCTGGACAAT	GAAGTGTTTT	TTGTTTTGTT
228241	GAGCAGTACT	CATCCTTTAT	TCTCACTCAC	CTCAAAACAT	TTCAATGGAG	TATTTTTTTG
228301	CTTTATGATC	TGGATGAGCC	CTTARARCAM	ACAGTAGCTG	AGAATTTATT	TCATAGTACT
228361	TTTGTTCCAC	ACTGTGGAGC	ATTCACARACAT	TGTAATATTA	ACTTAGCTGG	GAACAGAAAT
228421	TCCAATATGA	AATTTGTCTT	TTAGGAACA	GTATTGACTT	CCTGCTAGTC	TCTTCTGATG
228481	GGTGCTATAG	GGAAGTCTAG	ATCCTCCAAM	ACTITITGTA	GGAGAGCTAT	GTTTAGGCTA
228541	CATTTTCTCT	GATTCTCTTT	CTGGTCGTCT	TECTTCACCA	AGATGTGCCA	AGGTGTTAAT
228601	TGATTGTCCT	TGCTTTTTGG	TCTTTACTA	TAGAGTTTCC	TICGATTTTG	TTTTATTTAG
228661	AACCTTGTTG	CAATTTGTTT	TCTTTTCTCC	GAATCTCTCT	TCTATTTATC	TGTATGGTAA
228721	TTTCTCCATG	CCCATCTTTC	TACTCCACCC	IGACTITCAT	TTTTGGACCT	TTTACTTTGC
228781	TTCCATCAAT	GACTTTTTGG TTCAACTTAT	TTCCTAAAAT	AGGCAAACAC	TTTCCAAAGT	CTTTCTCAAT
228841	CCCTCCTTCC	ACTTTAGAAA	CCAAACCCAT	CCL CL COURTS	TGTGCCTATG	TCCACAATAT
228901	TGTAAACACT	TTCTGGTTGT	CAACAAGGCAI	CCACACTTTA	TTTAGGTGCA	ATGCCTGAAG
228961	TGCTAATGAT	TAACACATTC	ACCTTCCCTC	TTCCTTCCAA	ATATTGGTTT	GGGGATAACC
229021	GCTGTGTGTA	TTTTTTTAA	TCACTCACAA	TATOGATA	TGCTCCCTCT	TCTTTTATCT
229081	AGAGGACTGG	CCAGAGTGGG	AATGTTCTCA	ATTCACARTA	ATTGTATGTT	TTATTATAAG
229141	GAACTCATTC	TTTCAAATGA	ACCTCCCATA	TTTTCCCACA	ACTGAAGCAG	TACAGGATAG
229201	TTTAAAAAAC	TTTCAAATGA TTGATATGAA	TCATACAATA	1111CCCAGA	GCACCAAATT	TCAATATATA
229261	TATGTCATGA	AATACTTATT	CTAATTATAC	TCL CTCTTC	AACTITTATT	AAAATAAACT
229321	TGTTTAATGT	TTTCTTTTAT	TTACAAAACA	ATTATION	TCTTATTTCA	TCTTATAACA
229381	CAAGTTAAAA	ATATTCAAAG	GAATGCCTAA	ATTIATTIT	TGATGAAAAG	TTTTAGAAAT
229441	ATCAAAAGAG	TCTGAAGACC	ATTTACCTAT	CCARATTCE	MATTETTTAC .	ATGTTGTACA
229501	TCTAATATTT	ACTATTTATA	ATCCTTANA	ATTTCCCTTA	TATTTTTAAG	CAGTATCCCT
229561	CCCAGGAGAC	GGAGGTTGCA	GTGAGCCAAC	ATTIGCCTIA	GCACAGGAGA ,	ATTGCTTGAA
229621	AGAGTGAGAC	TCTGTCTCAA	AAAAAAAAA	ALAGIGCCAC	AAAAAAGGGG	LCTCGGCGAC
229681	AAACAAACAA	AAAAATCCGC	CTTAACATTA	TTTCTTCXT	AAAAAAGGCC A	HAAAACAAAT
229741	CTAGTTTCCC	TTTCCTCTCA	GCCCATTGTC	ATATTTTCALL	MAAAACTTTC	TAATACTA
229801	GACATATGAG	GTTTTTGTTT	Thirting	TTGGACATGG	ACTOROGET :	IGCTTTGTAG
229861	GCTGGAGTGC	AATGGCGCAA	TCTTGGCTCA	CTCCAACCTC	AGICTCCCTC '	rgTTGCCCGT
229921	TTCTCCTGCC	TCAGCCTTCC	AAGTAGCTGG	GATTACAGGG	ACCCACTAGE	SI I'CAAGCAA
229981	AATTTTTGTA	TTTCTGGTAG	AGACGGGGTT	TCACCATGTT	GGCCACIACC /	ACGCC TGGCT
				ACCAIGII	GGCCMGGCIG (JICICGAACT

Figure 9 (Pag 71 of 74)

230041	CCTGACCTC	A AGTGATCCAC	AATCCTTGG	CTCCCAAAG	GCTATGATTA	G11G01G1
230101		- MOCCAGMAIN	L TATGTTCAT	「		
230161		" GITWCITICI	IGAGAAAAT	י דרדנאאארו	**************************************	
230221			CATATTGAGI	محمد سلمساسات المساور ال		
230281		· CIGINGIIMA	L AGAAACCAC	· TOTOTOTOTO		
230341		· TIGNOGIGG	GITACTCTG	L GAATCAAAGC	* *********	
230401		S GICTINGCIG	I AGCAACTTG	CTCCATTCT	מתת תמידות הכיי	
230461	TGTATTTTCC	CTCTACTCAA	CATTTAAGGT	CTCAGAAGAT	AATATAATTG	GGCTTGAACT
230521	AGTAAAGTGC	CTCACTCTTTT	GCTTTAACAA	ACCCTAGAGA	GCTGGTAGGC	GTGAAATTTA
230581	CAGACCGTTT	TAGCTTCCAA	AGGGAGTTCA	GGACACCATO	ATTCACGACC	AGAGCCTCAA
230641	ACACATAATT	GAGAAAAGAT	AGTTCCACCA	AATAAAATT	AAATGCTGAC	ACAATACATC
230701	AAGAAATCTT	GGAAATAGGT	TTATATAAAA		CCTTTTTTAT	AAGAAGGGGT
230761	TAGGACCAGT	TCTACTTAAG	CCACCCATTT	GCCDDDDTDD	AGTGAGAATC	TGTTATGGAA
230821	GGGACTCCTC	TTTGTAGCTC	CAAGTGCCAC	TALL MOODES	TAGGACCTGA	GTTTCTTTTG
230881	AGGTGATTTC	AGTTAATATG	ATCAATTATT	ייא א מידידער בייני די א א מידידער מידי	GGCTCTAATG	GCTATAAGCC
230941	ACGGAGCCCA	TCAGCATTCC	CTGCAGGGAA	CTGCAGTGGG	TTTTATCAAC	TGCAGAGGGA
231001	AGCTTTCAAC	TGTTTTGAAA	TCACTTTCAG	GGTGGTCATC	TAGTTGCTTT	TTGAACAGCT
231061	GAAGATGATT	CTGCCTCTTT	TAATATGTGA	CTCCTCACAT	TCAGAAAGTG	TTTGAAATCA
231121	TTAAGAGTGA	ATTACCCTCA	GTGGTCCAGC	CTCCTCAGAI	CCACATCTAA	CTCGCTAGTC
231181	TGGGGGAACT	ATCAGAGAAA	TTGGTGCCAT	GCACATARCA	GGAAGGCACA	CCCTATCCCC
231241	GAGCCCCGCA	TGATGAAAAT	CAGTGGACAG	CATCATTA	TACAACTTTG	GTGAAGCAGA
231301	GGAGCATGAA	AATCCAGGCC	AATCTGGCAC	CATCALIATI	AATTTTTGTT	TAATCACCCA
231361	GAACCGATTC	TGATGAATGA	CTGTTTAGCC	ATTTTACACT	GTGGCATACG	GGAGTTCTTG
231421	CATACAGAGG	TTGGATGTAA	ACGGGCCTTT	CCCCTCTCTT	ATGAACATAG	TGGCTGCTGG
231481	AACTGTGTCA	CATAGGTTCC	AAATGGTGGC	CTGAATACTA	TTTACAACTA	ACAGGAACTA
231541	AAATTGAGTA	AGTCTTTTCC	TCTTTTCCAG	ATACCATCAT	TATTCATATA	AGGTACAATG
231601	GTTAACTATT	TGTATTTGGT	AATTTTTAAT	ACANATOTA	TATTCATATA	TTTCTTCAAA
231661	TCTTTAGTCT	TAAGGTTGAT	GCTCTCCATG	TCCTTCCNNN	AAAAGGTATG	TCAAGTTTAG
231721	TATATCCTCG	CCTTCAGATG	GGATTATTCC	ATTTTCTAA	TTGTTAATAT	TTGCTTTTAT
231781	CCACTTTTTT	TGTGGCTCTG	GGTGAGATGC	TATACCTACA	ATGACAAGTG	ATACTTTGAG
231841	TGTCCCTGTC	ACAAAAGTGG	ATAGCCTAAG	TCCTCACTCT	TACCTCCACT	ATACGTGTGT
231901	GTATCACACA	CCAGCCGTAT	GCCAGGCACC	ACTCTACCTC	CTAGGGATAC	CCAAATATAT
231961	AGACAAATGC	AACCCCTGCC	CATGTGAAAG	ACTUTAGGIG	AATAAATAAG	AGCAGTAAAC
232021	GTTATATGGA	GGTGGCAAAT	GCTAAAAAGA	ADAMIANGAC	AGGCAAGAGG	TAAAGTGCAT
232081	AAGATGACAT	TTGGGTAAAA	GCCCATGTAT	ATATCTTCTA	TIGGTTTTAT	ACTCATTGAA
232141	AGCCCTGACT	AATACACAAT	GACTTTGAGA	AGTTACTCCC	TTTTGATTTA	TTCTCTGGAG
232201	CGGAGTGCTG	AGAGCCTTCT	TAGTGTGTAT	TCAGTGTTTT	AAGAGAGCTT	TCACACTATT
232261	AATAAATAGG	ACAAAATTTA	TCCAAACTTA	AGCCTTGCTT	TAGGTAAAAG	GTGGATGAAT
232321	ACAAGGTAGA	AGGTTATTAT	TTGACATTTA	AATCCAACTC	AAGACTAATA	GGCTCCTCTT
232381	ATTAAAAGTT	TTTAAATCAC	AACTGCGTGC	AAAATAAATG	GAACTGCCAT	AGACTAATTA
232441	TGTGCATGAG	TGGTGTGCAT	GGGAGACAGC	ACGAAGCTAA	TCCCACTCAT	GCTCGCCAAG
232501	GCTCCATTTT	TCTCCTAAAA	TCAGTAAGAC	AGAAGCTAA	CAGATTATCA	CTTGCAGGTT
232561	TTAAACACAG	CAGTAGCATT	TGGAAGGGGT	TCCTCTCATT	AGGCAGTGCC	AGAGCCCTAG
232621	AAGAGATGAA	CAAGCCCTGT	ATCTGAAGCC	ATCATGCCTA	GTTATGGTCC	IGACCACAAC
232681	ATGATGCCTG	GAAGGGAGGC	CCCCTGCACC	CTAGAAACCT	GGGTGGGTTC	CCGACTGTTC
232741	TTTACTGCTA	AAAACCCTCT	TCTTTGGATC	TGGACTTTAC	CTCTATCTGA	TACTGTCTGC
232801	TAATATATGA	TTTGGCACTG	AGTCTGTCAC	TGCTGCTAAC	TCAGCAGTTC	TTTTTTTC
232861	GCCCCATTGC	CTCACAGAAA	GAATTTCATA	COTTCCACCA	TCAGCAGTTC T	TAGGGTCATT
232921	TTTGATTTCA	GCATTGCTAT	TTTTTTCTC	GGGTGTTCCA	GCTCTCTCTCC ?	TCATTATAC
232981	GTCTTGTTGG	TTTTCTGCTA	ACTCCTGCTT	The second of th	TTTTTTTTT I	CCTTCCCAT
233041	TCGTTCTGTC	ACCCAGGCTG	GAGTGCAGTG	GCACAATCTC	GGCTCACTGC A	AGACGGAGTC
233101	TCCCGGGTTC	AAGCTATTCT	CCTGCCTCAG	CCTCCCAAGT	AGCTGGGACT	ACCTUCGCC
233161	ACCACTATGC	CCCACTAATT	TTTGTATTTT	TAGTATTGCT	GTCATCAATC	ACAGGCGCTC
233221	GAAGCACCTA	GAAACTCTAA	TTCTTTGTAG	GTATCAAACC	TAGGACTCT T	ACATGTCCA
		-			- *WORWCICI, J	ICCICTAAT

Figure 9 (Page 72 of 74)

233281	СОСОВТОТОТ	AATCCCTGAT	TCCCDDDCDC	CCTCTTTTCA	татасаттт	ССУСТСТАСА
233341		CCTGGAAAGC	-		-	
233401		AAGAATCAGT	_			
233461		TATCTGAATG				
233521		TTGCTGCCCA				
233581		GGTTCAAGTG				
233641		CACACCTGGC				
233701		TTCCTCGAAC				
233761	· · · · · ·	ACAGGTGTGA				
233821		AGTCTGACAC				
233881		CAGATTTCCT				
233941		GCCTCAAGTT				
234001		CTGAAACAGG				
234061		CATTTTTCCT				
234121		TGTTAATTTA				
234181		TTAAAAAATT				
234241		AAATGAGGAA				
234301		TGTTAGCTTT				
234361		ATGCTCATAA				
234421		AATCCCAGCG				
234481	TTTGAGATCA	GCCTGGGCAA	CATAGTGAGA	CCCTGCCTCT	GTAGAAATAA	ACAAAAATTA
234541	GCTGGATATG	GTGGTGCATG	CTTGTACTCC	TAGCTACTTG	GGAGGTTGAG	GCAGGAGGAT
234601	CCTTTGAGTC	CAGGAGTTTG	AGGCTGCAGT	GAGCTATAAT	CACCCACTGC	ACTATAGCAT
234661		GTGAGAACTT				
234721	TATAAACAAA	ACTITIGITT	CAAAATATGT	AATATTTAGC	ACTAAAGAAT	TCTGAATTGT
234781		AGTACTTAAA				
234841	AGTATAATTT	TTATCCAGAA	AATCATCCAT	ATCAGCAAGC	TAAACTTTCT	CAAAATGACA
234901	TATCCATGTA	ATTAGCTCCC	AGGTAATTAG	CAGGCAGCCT	CTACTCAGGT	TGAGTATTCC
234961	TAATCTAAAA	ATTGGAAATT	CAAAATGCTC	CAAAATCTGC	AACTTTTTGA	ATGCTAACAT
235021	GATTCTCAAA	GGAGTGCTCA	TGGAGTATTT	CAGATTTTGG	ATTTTTGGAT	TTGAGATACT
235081	CAGTATAATG	CAAACATTCC	AAATCTGAAA	AAATCTGAAA	TACTTCTGGT	TCTAAGCATA
235141	AGGGATACTC	AACGTGTGTT	AGCTAATTAG	ACCCTTCATG	GTCTCTTCTA	GACCTCAGCT
235201	TCTTCAAGGT	AACCTCTATC	CTCACTTCTA	ATAGCATGAA	CTTTTCTGTT	TTAGAATAAT
235261	TTGGATTTTC	AGGAAAGTTG	CAAAGATAGT	ACAAAGACAG	TACAGGAGAG	TTCCCATATA
235321	TCTTTCACCT	AGCTTTCCCC	CATTGTTAGG	ATTTTACATT	ATTATGATAC	ATTTGTCAAA
235381	TATAAGCAAC	TCACATTGAT	ACATGAAACT	CTATTAACCA	AACCCTAGAC	TTTATGTGGA
235441	TTTCACCACT	GTTTCCACTA	ATGTTTTCTT	TCTGTTCCAA	GGTCCAATCT	GGAATACCAC
235501	ACTGCATTTT	CTTGTCATAT	CTCCCTAGTC	TTTTTTTGTC	TGTGACAATG	TCTCAGTCTT
235561		TCATGACCTT				
235621		AGATTTTTTG				
235681		GAACATGATA				
235741		TTCAGGTTTC				
235801		AATTTTGTTT				
235861		AAATCTTAGA				
235921		ACATTCTAAA				
235981		ACTTTTGTAT				
236041		CCAGTACTAT				
236101	CTTAGTTGGC	AATATTTTTG	TIGGTTTATI	TCTAGACTGT	TTATCTCATT	CCACTGATTT
236161	GTGTCTATCT	TTTTGACAAA	ACTGTTGATT	ACAGTAAGCT	TTGAAATAGT	TCATTTTTTG
236221	TGTCAACTTG	ACTGAGTCAG	GGGATAACCA	GCTATCTGGT	TAAACATTAT	TTCTGGCTGT
236281	GTTTGTGAGC	GTGTTTCTGG	ATGAGATTAG	CCTTTGAATA	GGTGATCCTA	GTAAAGTAAA
236341	CTGTCTTTCC	CAGTGTGGAT	GGCATTATGC	CACCTGATAT	TCAGGGTCTG	AATAGAAGAA
236401	AAGGCAGAGG	AAGGGGGAAT	TTGGGCCTTT	TTTTCTGCCT	CACTGCTTGA	GCTGGGACAT
236461	CTCATCTGGT	CTCCTGCTCT	TGAACTGGG	TTTACATCAT	CAGTTCCTCT	GGTTCTCAGG
230701	CICAICIOGI					_

Figure 9 (Page 73 of 74)

236521	CCTTCAGATT	CAGACTGAAT	CATACCACCA	GCTTTCCTGG	GTCTCCAGCT	TGCAGATTAC
236581	AGATCATGGG	ACTCCTCATC	TTCCATAAAT	GCATGAGCCA	ATTCACTOCA	TGTCCTTGAA
236641	AACTGCCCCA	CTGCAGATTA	VCCCunnama	CCACTAGGTG	ATTCAGTCTA	TGTCCTTGAA
236701	CAGATTTCCC	TTONTONIA	AGGCTTTTT	CCACTAGGTG	AAATAAAGAA	GCTTGTTAGA
236761	DCTCCC C	TICATCCAGT	GCCCTCTCCT	CTTTAAGTTA	CAACACATTG	GCTACACCTA
	AGIGCAGGGG	TGGGGATGAG	GGTATAGTCC	TCTTGTTTGC	TGAGAAGAGA	ACTGTATTCG
236821	GAMAGCTCTA	GAAGTGTTTG	ATACATACAT	AAACAAGGCA	TGGTTTTTGC	ACTTA ATTEC
236881	ACATTACATT	TTTCCCAGAA	AAAAAGGAAT	GTATAGGCAT	CACCTAACTC	TOTAL TIC
236941	AGTCATTCTT	CCTGATTATC	D D D C C C D D D D C	ACCOUNT AGE TO	CACGIAACIG	TACTAGCTGG
237001	ACA ACTA CTT	TTTCC	AAAGGIAAAC	AGTTATTAAT	CCTATACCAA	GATGTCAAGG
237061	AGAAGIACII	TIGGAACACA	AGGAATTCTC	TGGGAGTCCT	TACTACTCTC	AAGCCCAGTG
	AAAAAGTTAA	TGAAAAACTA	TAGTACCTTC	CTATAAGCTG	GATGACTAAT	TACCAGGGTC
237121	ATTTAGGAAT	TTGCCTTACC	AAGTAAAACA	TAAGGGCAGC	TGAGGTGCTG	ACTCAACACA
237181	AATGGAGCAT	AGAATAAGAG	TAGTAAAGAA	TGCCAAAAAT	CCTCTCATCT	ACIGAAGACA
237241	AAAAGGAGCT	ATAAAGCCTT	TACCTATOTO	CACACACACA	GCIGICAIGT	ATCCATTGAC
237301	TGTGTGTGTGTG	TOTOTOTOTO	TAGGIATITT	CACACTTGCT	CTGTTACGTA	AATGTATGTG
~3.301	1919191919	TGTGTGTGTG	TGTGTG			

Figure 9 (Page 74 of 74)

International application No. PCT/US97/17658

1	to International Patent Classification (IPC) or to both national classification and IPC	·-··
	LDS SEARCHED	
I	documentation searched (classification system followed by classification symbols)	
U.S. :	536/23.5; 435/6, 70.1, 325, 320.1	
Documenta	tion searched other than minimum documentation to the extent that such documents are included	in the fields search
<u> </u>		
1	data base consulted during the international search (name of data base and, where practicable	e, search terms use
	ALOG'S BIOTECH cluster. omatosis, BTF1, BTF2, BTF3, BTF4, NTP-3, NTP-4, RoRet, butyrophilin, type 1 sodium t	rananad
C. DOC	CUMENTS CONSIDERED TO BE RELEVANT	
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	hemochromatosis locus. Genome Research. May 1997, Vol. 7, No.	- .
	5, pages 441-456, see entire document.	
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^	FISCHER, L. et al. Cloning of the 62-kilodalton component of basic transcription factor BTF2. Science. 04 September 1992, Vol.	28-33, 71
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	25 January 1991, Vol. 251, pages 424-426, see entire document.	
X Furth	er documents are listed in the continuation of Box C. See patent family annex.	
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International application No. PCT/US97/17658

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C (Continue	ation). DOCUMENTS CONSIDERED TO BE RELEVANT		
Category*	Citation of document, with indication, where appropriate, of the releva	int passages	Relevant to claim N
x	ZHENG, X.M. et al. Sequencing and expression of com DNA for the general transcription factor BTF3. Nature. 1990, Vol. 344, pages 556-559, see entire document.	-1	34-39, 72
	PANTEGHINI, M. Electrophoretic fractionation of 5'-nu Clinical Chemistry. February 1994, Vol. 40, No. 2, pages see entire document.	ucleotidase. s 190-196,	52-57, 75
1	BURT, M. J. et al. A 4.5-megabase YAC Contig and ph map over the hemochromatosis gene region. Genomics. 1 1996, Vol. 33, No. 2, pages 153-158, see entire documen	5 April	1-6 7-20, 22-77
li	VERNET, C. et al. Evolutionary study of multigenic families mapping close to the human MHC Class I region. J. Mol. Evol. November 1993, Vol. 37, No. 6, pages 600-612, see abstract in particular.		1-20, 22-77
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Form PCT/ISA/210 (continuation of second sheet)(July 1992)*

International application No. PCT/US97/17658

Box 1 Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)
This international report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:
t. Claims Nos.: because they relate to subject matter not required to be searched by this Authority, namely:
2. Claims Nos.: because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:
Claims Nos.: because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).
Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)
This International Searching Authority found multiple inventions in this international application, as follows:
Please See Extra Sheet.
As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims.
As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3. As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos.:
4. No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claims Nos
Remark on Protest The additional search fees were accompanied by the applicant's protest
X No protest accompanied the payment of additional search fees

Form PCT/ISA/210 (continuation of first sheet(1))(July 1992) *

International application No. PCT/US97/17658

BOX II. OBSERVATIONS WHERE UNITY OF INVENTION WAS LACKING This ISA found multiple inventions as follows:

This application contains the following inventions or groups of inventions which are not so linked as to form a single inventive concept under PCT Rule 13.1. In order for all inventions to be searched, the appropriate additional search fees must be paid.

Group I, claim(s)1-20, drawn to polynucleotide sequences containing at least one polymorphic site, polypeptides encoded thereby, antibodies to said polypeptides and a method to determine the presence of the HFE gene mutation.

Group II, claim 21, drawn to the lymphoblastoid line atcc crl-12371.

Group III, claim(s) 22-27 and 70, drawn to BTF1 nucleic acids, gene products, vectors and antibodies.

Group IV, claim(s)28-33 and 71, drawn to BTF2 nucleic acids, gene products, vectors and antibodies.

Group V, claim(s) 34-39 and 72, drawn to BTF3 nucleic acids, gene products, vectors and antibodies.

Group VI, claim(s) 40-45 and 73, drawn to BTF4 nucleic acids, gene products, vectors and antibodies.

Group VII, claim(s) 46-51 and 74, drawn to BTP5 nucleic acids, gene products, vectors and antibodies.

Group VIII, claim(s) 52-57 and 75, drawn to NPT3 nucleic acids, gene products, vectors and antibodies.

Group IX, claim(s) 58-63 and 76, drawn to NPT4 nucleic acids, gene products, vectors and antibodies.

Group X, claim(s) 64-69 and 77, drawn to RoRet nucleic acids, gene products, vectors and antibodies.

The inventions listed as Groups I-X do not relate to a single inventive concept under PCT Rule 13.1 because, under PCT Rule 13.2, they lack the same or corresponding special technical features for the following reasons: Groups I and III-X are drawn to physically different genes and their gene products and each therefore constitutes a separate invention. The lymphoblastoid cell line of Group II is not dependent upon the vectors of any of the Groups I and III-X and therefore constitutes a separate invention. Accordingly, the claims are not so linked by a special technical feature within the meaning of PCT Rule 13.2 so as to form a single inventive concept.

Form PCT/ISA/210 (extra sheet)(July 1992)*

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